

Automated Alzheimer Disease Detection Model by inducing an Efficient Fusion Strategy for Image-based Classification

Arun Kuchanur¹ Dr.Kirankumari Patil²

^{1,2}Department of Computer Science & Engineering

^{1,2}REVA ITM Visvesvaraya Technological University

Abstract— This paper presents a new fully automated image based classification method to evaluate the neurodegenerative disease, by feature extraction from brain magnetic resonance (MR) images based on support vector machine (SVM). Neurodegenerative diseases involves large variety of mental disorders, the evaluation of disease is not particularly related to visual differences carried out by radiologists. Such analysis that may examines the disease, which may not give comprehensive results to disease. Hence by this paper we introduced general and special visualization software designed in Visual Basics 2010. Application will produce quantitative and clinical analysis of MR images of brain. We use the Alzheimer's disease (AD) as the case study; the results are affected regions of AD, and a plotted graph for regions of interest and percentage of death of neurons. This is achieved by fusion strategy, which combines bottom-up information flow and top-down information flow. Bottom-up includes Multiscale analysis of features from different images, where as top-down includes learning phase and fusion problem. Finally difference in individuality of regions found by this approach is highly correlated to clinical studies of Alzheimer's disease.

Key words: Support vector Machines (Svms), Alzheimer's Disease (AD), Automated Pattern Recognition, Computer-Assisted Image Analysis, Resonance Imaging (MRI)

I. INTRODUCTION

AD is first reported by the neurologist Alois Alzheimer from German, is a disease makes a difference in the physical part of the brain. Death of neurons in the brain is the main cause of the disease, in turn which results in memory loss and failing in strength of the mental action or gradual slant down of the acquiring knowledge. A neuro-degenerative type of disease, it starts slowly and grows increasingly worse. AD is the usual reason of loss of memory and mood changes, i.e. dementia reports a group of indications which also involves problems with communication and reasoning. These indications happen whenever the brain cells are destructed by definite diseases and conditions-in case of death of neurons or including AD. There might be shortfall of significant chemicals within the brain may cause certain diseases. Such chemicals will be participating in the exchange of messages within the brain. Alzheimer's disease-one of the progressively developing disease; means disease moderately increases in stages or increases day by day, and then other parts of the brain may also get damaged. If this occurs, the indications become more serious. Initially these indications are often neglected for age-related issues or instance of stress. The short term memory loss is most common initial indication of AD—difficulty in recalling or remembering recent activities. The identification of symptoms is normally complying with the examinations of the person thinking capability and behaviour, usually come

after by a brain scan if obtained; however, a detailed inspection of brain tissue is required for a proper conclusion. As the disease gradually increases, symptoms can also involve uncertainty, irritability, violent behaviour, sudden mood changes, trouble with communication, and short or long term memory loss.

At analyzing structural brain MR images, an important thing is to find changes in anatomical structure, either local or global, changes may relate to functional disturbances or interruptions. In particular, radiologists evaluate MR images by looking at particular regions and then compare them by finding differences [1].

In the computational attempt of emulating the human vision process—a synchronized collaborative work between the brain and low level visual mechanisms—the concept of visual attention has introduced a generation of techniques that are able to transform an image into a hierarchy of relevant regions, known as salient regions. Relevant regions in radiological terms may be defined as those image areas that are visually altered and are entailed with a certain degree of clinical interpretability. Nevertheless, most methods used to compare brains establish local rather than regional (salient) differences.

Currently, a morphometric brain analysis consists of a set of strategies aimed to extract and quantify anatomical differences between groups of subjects. Commonly, this analysis comprises two main processes: first, all images are warped or registered together to a common reference frame or template, and second, a quantification of the estimated local deformation required to register is computed, producing specific measurements of interest. voxel-based morphometry (VBM) [2] and deformation-based morphometry (DBM) [3] are currently the most used techniques to compare populations. In VBM, local differences, found in brain tissue segmentations, are voxel-by-voxel statistically analyzed, while DBM statistically compares in-formation coming from the deformations fields obtained after registration to the template. With these methods, one-to-one correspondences between subjects are assumed and statistics are computed for the same voxel across all subjects. However, conclusions are limited when the same structure may be partially present, or when a single anatomical region may exhibit multiple shapes across the population. On the other hand, some pathologies may affect not only a unique anatomical structure or even contiguous regions, but localized structures separated from each other. These kinds of patterns are difficult to find and analyze with these classic morphometric techniques. A recent proposal, the feature -based morphometry (FBM) [4], copes with these issues by modeling the image as a collage of local scale-invariant features and by learning, from them, a probabilistic model that reflects group- related anatomical characteristics. However, these approaches disregard the local statistical dependences, and then subtle changes are hardly detected;

exactly the opposite strategy used by the radiologists, who analyze regions rather than pixels [5].

II. LITERATURE SURVEY

Existing studies suggest neuroimaging may become a valuable tool in the early diagnosis of neurodegenerative diseases by extracting anatomical patterns and revealing hidden relations from structural magnetic resonance (MR) images. The value of neuroimaging against clinical, neuropsychological and biochemical analysis remains to be demonstrated in large representative populations, yet there exists sufficient evidence in small series of patients with different states of neurodegenerative disorders. The process that an expert follows when examining a particular case involves two different kinds of tasks: those related with image perception, such as visual search or exploration paths, and others associated with cognitive skills, mainly related to diagnostic reasoning and decision making [1].

According to author [3] 1998, their point was to show a system for identifying gross plainly visible anatomical varieties in the cerebrum of different subjects. The procedure included spatially covering the basic MR outputs of distinctive subjects so that they all adjust to the same stereotypic arrangement. Multivariate insights were utilized to portray the evaluated nonlinear disfigurements that happen. To show the procedure they compared the gross morphometry of brains of both male and female subjects. They additionally evaluated mind asymmetry, the impact of handedness and relations among these Effects.

According to author [5] in 2009, they had inquired about component extraction procedure for the determination of Alzheimer's ailment on MRI pictures of the mind through voxel based morphometry (VBM). The gatherings of voxel areas recognized by the VBM were utilized to pick the voxel force variables on which the characterization qualities were processed. They investigated the advantages of the data from the first MRI registries and GM division volumes. In this exploration they additionally connected the bolster vector machine calculation to characterize patients mellow Alzheimer malady versus control. The study had been done on MRI sweeps of 98 females, after fastidious demographic race from open access arrangement of imaging studies (OASIS) database, which was a bigger number contrasted with current archived studies.

In their work they had scrutinized element extraction procedures grounded on VBM examination to sort MRI volumes of patients with Alzheimer's illness and controls. They had recognized distinctive plans for the SPM of the VBM and they found that the fundamental GLM outline outputs covariates can distinguish unpretentious changes in AD patients and typical subjects that prompt organizing of SVM more tasteful with a differential precision of 87.5 %. They looked at their results on a littler populace of AD patients and controls to the ones achieved with a standard VBM examine utilizing single group and ascertained an order precision of 63.3% by means of cross acceptance. Thus the outcomes revealed in this theory alongside the exploratory philosophy utilized could be used by the neuroscience group exploring on the AD. The further studies may incorporate extraction of characters in light of other morphometric methods specifically disfigurement based morphometry.

According to author [7] 2012, they concocted a novel PC helped determination (CAD) technique for right on time and simple conclusion of AD sickness on premise of non-negative factorisation ((NMF) and SVM (bolster vector machines) with jumps of certainty. The CAD instrument is modified for the study and categorisation of practical mind examines. For this system two distinctive mind checks databases are picked: a solitary photon outflow processed tomography (SPECT) database and (PET) positron emanation tomography pictures, the two bearing information for both AD patients and ordinary sound controls as a source. These reports are measured by utilizing the fisher discriminate proportion (FDR) and (NMF) non negative framework factorisation for trait determination and disclosure of the most proper components. The acquired NMF changed arrangements of data which contain a substantiated number of components, are gathered with a SVM based classifier with jumps of certainty for choice. The set forth NMF – SVM technique gains up to 91% order exactness with high affectability and specificity rates (more than 90%) . this NMF – SVM CAD method turns into an exact strategy for SPECT PET AD scans.

III. PROPOSED SYSTEM

This proposed approach is an automated image based analysis method, Moreover excellent approach then visual analysis made by the radiologist. In our proposed method system takes MR images as the inputs and gives the output as normal or pathological conditions of the brain. The model is based on extraction of saliency maps and which also includes learning phase, which shows the slight alterations of visual description of radiologist. The method also does the analysis of salient maps at different scales that are combined optimally. The method uses learning process to classify and detect disease, so some patterns of brain are previously learnt and are related to normal or pathological conditions, and then input MR images are compared with learned visual patterns. This is not about interested point in the brain structure but about salient regions, thereby the complete brain structure is classified as normal or diseased one.

A main contribution of this work is a fusion strategy that learns, from training data, the discriminant structural patterns of neurological disorders, in particular, the Alzheimer's disease. Another important contribution is the model interpretability since the learned patterns can be mapped to the original brain and used to quantitatively estimate the importance of each region, normal or pathological, for the final classification. Technical contributions include: the use of a 3-D multi-scale analysis of the brain saliency inspired by what radiologists do when examining cases, the use of low-level features that sparsify data, the formulation of a model and fusion strategies as a max-margin multiple-kernel optimization problem, and a regional analysis method that completely avoids any non-rigid preregistration step, which at the end constitutes another important variability source. An extensive parameter analysis of the influence of the image features as discriminative factors is also carried out. The classification accuracy between normal controls and probable AD subjects is improved by applying this approach, outperforming a recently proposed technique (FBM).

To the best of our knowledge, this kind of visual-saliency-based pattern extraction approach has not been previously investigated for AD characterization and classification in structural MR images.

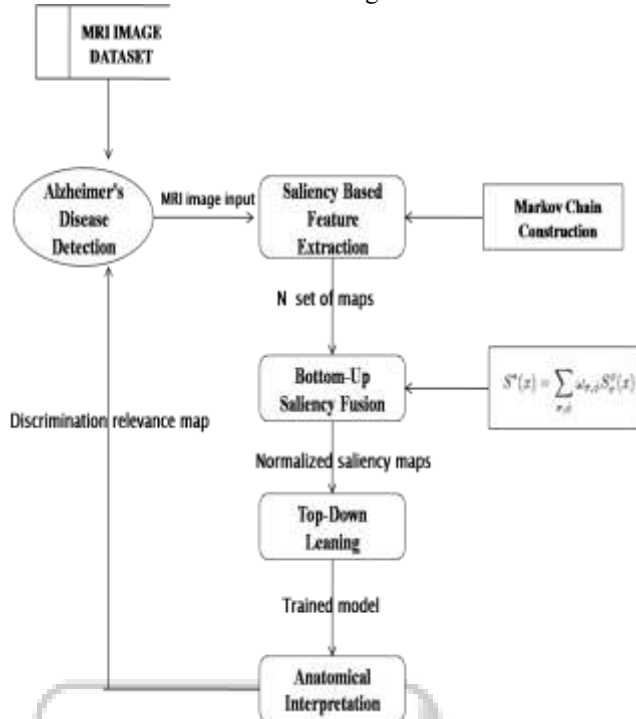


Fig 1: Overview of the proposed method. Different feature-scale saliency maps are extracted, then a learning algorithm fuses optimally this information to feed a SVM classifier, to produce both a classification model as well as maps of relevant anatomical regions.

IV. IMPLEMENTATION

A. Calculation of saliency maps:

The MR Images of structural brains are used to search of individual patterns among all the visual (“anatomical”) area of brain.

Here in this module, evaluation of saliency region is starts by absorbing a group of features from the brain volume “x”, i.e input image. Module implementation Procedure is given below. Steps:1: perform multiscale decomposition on structural brain volume “x”. 2: Extraction of feature per each scale. 3: Draw fully connected graph for feature and scale combination. 4: Construction of Markov chain on graph, calculation of equilibrium distribution of Markov chain. 5: Normalize the evaluated MR image. 6: Returns the N sets of salient maps for each combination of feature and scale.

B. Kernel Level Extraction of Maps:

Here in this module, this evaluates the similarities between sets of salient regions found in previous module.

Kernel function: --

$K: X * X \rightarrow R$

$\Psi: X \rightarrow F$

$X \rightarrow$ brain volume,

$F \rightarrow$ feature space,

$R \rightarrow$ extracted region from each salient maps

Hence the kernel function may look like it measures the similar space between input feature spaces, i.e.

saliency maps. Module implementation Procedure is given below.

Steps 1: Normalize all saliency maps and calculate kernel level function for all feature–scale combination. 2: Prepare kernel matrices for each feature–scale combination. 3: Feature–scale kernel matrices are the output of this module.

C. Trained classification model:

Here in this module, construction of trained classification model will be carried out. Goal of the model is which uses the parameters from the previous kernel function and most individuality from images are represented in different saliency region maps. And uses the set of labeled training volumes as domain knowledge, training labels are codified as train classification model. Module implementation Procedure is given below.

Steps: 1: For each kernel matrices got from previous modules use N set of labels to solve the min max margin-discrimination. 2: Obtain the feature-scale kernel weighting vector. 3: Generate the optimal kernel using a linear combination of feature-scale kernel and the learned weights. 4: Solve the min max margin-discrimination to get single optimal kernel to obtain the classification model.

D. Anatomical analysis (SVM Classifier):

In this module, method determines the anatomical areas, which are related to the specific patterns and highlighted by yellow colour. The module actually does the quantification estimation of the extracted brain differences. There by it achieves clinical analysis for Alzheimer’s disease (AD). Module implementation Procedure is given below.

Steps:1: Module will consider the trained classification model and N sets of saliency maps as input. 2: Constructing master saliency maps using learned feature-scale kernel weighting vector from previous module. 3: Using trained classification model, construct an overall discrimination relevance map using a liner combination of the master saliency maps and their corresponding classification coefficients. 4: Align the discrimination relevance map to extract maximum relevance values of specific anatomical regions. 5: Returns discrimination relevance region R by yellow colour (pathological regions).

V. RESULTS

Below shown pictures are the experimental results which are taken from the .net framework

A. Step 1: MR image used as input to the morphometric analysis model



Fig. 2: Step 1

B. Step 2: Outline of the MR image.

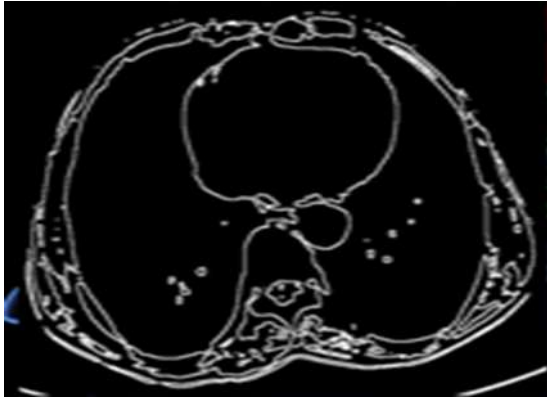


Fig. 3: Step 1

C. Step 3: kernel level abstraction of region.

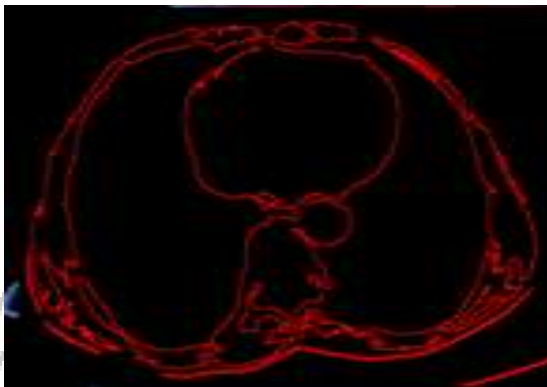


Fig. 4: Step 1

D. Step 4: Showing the affected area with yellow colour and numbering the regions.

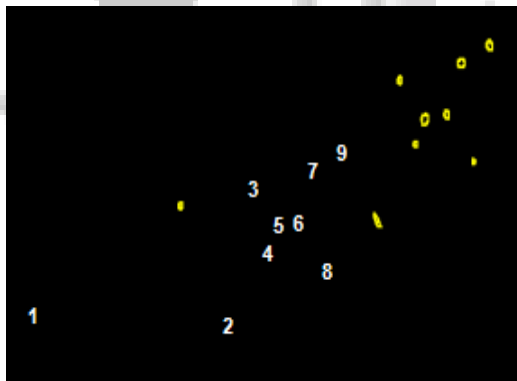


Fig. 5: Step 1

E. Step 5: The below graph represent the number of cell defected due AD.

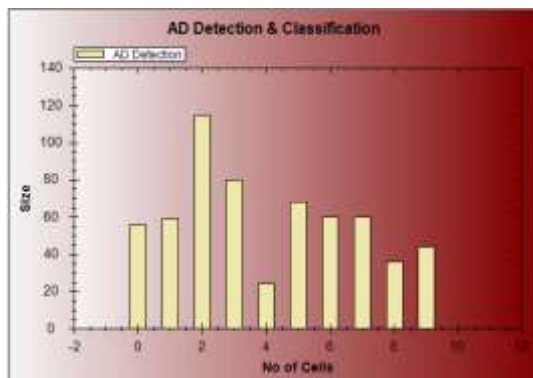


Fig. 6: Step 1

VI. CONCLUSION

The proposed approach is a completely programmed procedure that uncovers basic brain examples related to the vicinity of the AD in an ADNI MRI dataset. The basic thought behind the proposition is that which is possible to locate the different examples analyzed by a specialist surgeon may find in comparable pictures. This is refined utilizing a combination methodology that combines bottom up and top-down data streams, accomplishing precise groupings of likely person's with AD. The Bottom-up includes Multiscale analysis of features from different images is taken by a visual saliency technique that naturally highlights significant districts connected with the Alzheimer's disease analysis. Then again, top-down includes learning phase and fusion problem, which permits to choose the changes in the important piece of the delineation, recognizing examples related to diseased stages. As specified in the Introduction, data originating from voxel intensities (VBM) or disfigurement fields (DBM) is usually utilized for factual ID of anatomical between-gathering differences. The entire methodology permits to find anatomical areas with clinical implying that can be quantitatively identified with the conclusion, and along these lines, may be suitable for a target graduation and comprehension of the diverse AD stages.

By this approach we have found that principle changes are situated in horizontal directions, a roundabout confirmation that progressions happen likely in arranged regions however not precisely situated in a specific area. From the clinical point of view, the proposed procedure takes after the visual examinations done by radiologists when examining medical pictures, permitting likewise a quantitative determination of the mind anatomical locales which are diverse between exploratory gatherings. With a satisfactory and thorough assessment in bigger data sets, containing adequate cases of the distinctive AD stages, this strategy can be additionally utilized as a second analytic feeling in the present clinical usage.

REFERENCES

- [1] J. Beutel, H. Kundel, and R. Van Metter, Handbook of Medical Imaging. Bellingham, WA: SPIE Press, 2000, vol. 1, Phys. Psychophys.
- [2] J. Ashburner and K. Friston, "Voxel-based morphometry: The methods," Neuroimage, vol. 11, no. 6, pp. 805–821, Jun. 2000.
- [3] J. Ashburner et al., "Identifying global anatomical differences: Deformation-based morphometry," Hum. Brain Mapp., vol. 6, no. 5–6, pp. 348–357, 1998.
- [4] M. Toews, W. Wells, D. Collins, and T. Arbel, "Feature-based morphometry: Discovering group-related anatomical patterns," NeuroImage, vol. 49, no. 3, pp. 2318–2327, Feb. 2010.
- [5] H. Kundel, C. Nodine, D. Thickman, and L. Toto, "Searching for lung nodules a comparison of human performance with random and systematic scanning models," Invest. Radiol., vol. 22, no. 5, pp. 417–422, May 1987.
- [6] G. Orrù, W. Pettersson-Yeo, A. Marquand, G. Sartre, and A. Mechelli, "Using support vector machine to identify imaging biomarkers of neurological and

- psychiatric disease: A critical review,” *Neurosci. Biobehav. Rev.*, vol. 36, no. 4, pp. 1140–1152, Apr. 2012.
- [7] P. Padilla, M. López, J. Górriz, J. Ramirez, D. Salas-Gonzalez, and I. Álvarez, “NMF-SVM based tool applied to functional brain images for the diagnosis of Alzheimer’s disease,” *IEEE Trans. Med. Imag.*, vol. 31, no. 2, pp. 207–216, Feb. 2012.
- [8] M. García-Sebastián, A. Savio, M. Graña, and J. Villanúa, “On the use of morphometry based features for Alzheimer’s disease detection on MRI,” in *Bio-Inspired Systems: Computational and Ambient Intelligence*, ser. Lecture Notes in Computer Science. Berlin, Germany: Springer, 2009, vol. 5517, pp. 957–964.
- [9] N. Doan, B. van Lew, B. Lelieveldt, M. van Buchem, J. Reiber, and J. Milles, “Deformation texture-based features for classification in Alzheimer’s disease,” *SPIE Med. Imag.*, 2013.
- [10] M. Liu, D. Zhang, P. Yap, and D. Shen, “Hierarchical ensemble of multi-level classifiers for diagnosis of Alzheimer’s disease,” in *Machine Learning in Medical Imaging*, ser. Lecture Notes in Computer Science. Berlin, Germany: Springer, 2012, vol. 7588, pp. 27–35.
- [11] E. Westman et al., “Combining MRI and CSF measures for classification of Alzheimer’s disease and prediction of mild cognitive impairment conversion,” *NeuroImage*, vol. 62, no. 1, pp. 229–238, Aug. 2012.
- [12] B. Magnin et al., “Support vector machine-based classification of Alzheimer’s disease from whole-brain anatomical MRI,” *Neuroradiology*, vol. 51, no. 2, pp. 73–83, Feb. 2009.

