

An Intensive Learning of Deep Learning in Alzheimer's Disease Prediction

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Abstract—The senior population is disproportionately affected by Alzheimer's disease (AD), making it the most common form of neurodegenerative disorders. It has an impact on patients' life, causing cognitive capacities such as consciousness, speech, actions, and problem-solving to deteriorate with time. Unfortunately, there is currently no cure for AD, and progress has been slow on finding a cure. However, the focus was on use of current medical information to diagnose the cognitive status of a patient, which in fact shows that a computer can replicate the clinical decision-making process of a physician. These rapid improvements have made it possible to collect enormous amounts of multifunctional neuroimaging data, which has led to the recent deployment of deep learning to the prompt identification and classification of Alzheimer's disease. Computational intelligence study into Alzheimer's disease continues to be in its infancy, but it is expanding rapidly as even more hybrid information sources are being incorporated, and as transparency is increased through the use of explicit strategies that integrate knowledge of specific behaviours and paths associated with the disease. This finding can be further understood by comparing multiple data-driven methodologies based on large-scale organization healthcare information from AD risk stratification, which may one day lead to improved selection of people at risk for AD in clinical laboratories or early identification of AD in clinical studies.

Keywords—Alzheimer's illness, Disease progression, Deep learning, Cognitive deficiency.

I. INTRODUCTION

Alzheimer's syndrome is the neurodevelopmental disorder characterized by brain atrophy (shrinkage) and cell death. It is a common type of dementia, and it is marked by a gradual loss of capacity of the individual to operate independently. Dementia is by far the most common kind. Early indicators of the condition include losing details about recent events or talks. With the progression of the disease, people living with Alzheimer's will develop serious memory impairments and become incapable of performing daily duties. Medicines may improve symptoms temporarily or slowly. These treatments can sometimes contribute to maximizing functioning and maintaining independence in people with Alzheimer's illness [2].

Alzheimer's disease patients and carers can benefit from a variety of programmes and services. Alzheimer's ailment, a degenerative brain disorder, currently has no effective treatments or cures. Infections resulting from significant brain losses — such as dehydration, starvation, or infection — result in death as the disease progresses. Alzheimer's disease is caused by an unknown factor. The functioning of brain proteins, on the other hand, triggers a cascade of

damaging events in brain cells, disrupting the brain's workings (neurons). Damaged neurons lose their connections and finally die [7].

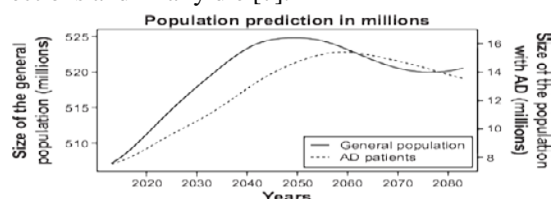


Fig. 1. Prediction of Population Vs AD patients

II. EASE OF USE

Genetic alterations cause Alzheimer's disease in less than 1% of cases, essentially guaranteeing that a person would get the condition. These unusual occurrences frequently result in mediaeval diseases. The damage usually begins in the memory-controlling area of the brain, although the initial symptoms do not occur for years. Neuron loss spreads to other parts of the brain in a predictable fashion. The brain had shrunk greatly during the late stages of the disease. Preclinical testing of Alzheimer's disease patients can lead to earlier detection of the disease and improved treatment options to delay its onset. The current AD biomarkers necessitate the acquisition of specimens or imaging data. E-health data, such as clinical records or organizational health data, on the other hand, do not require any additional time or effort to obtain. Furthermore, with the advent of digitization, the amount of such data has exploded [10].

Primary healthcare characteristics (maturity level, sexual orientation, profession), culture (physical exercise), midlife health-related risk factors (modest hypertension, body mass index, and total cholesterol), and cognitive assessments are often utilized to determine Alzheimer's illnesses. In clinical settings, evaluating whether these basic models, which are based on a small number of variables, can adequately account for the numerous etiologies of multifactor AD is a major challenge. In actuality, multi-factor modelling predicts the best disease risk, according to a meta-analysis research, however single-factor modelling does not reveal that successful AD risk prediction demands a wide range of factors. Here the magnitude is tested to which a data driven deep learning framework collects significant data from huge healthcare data that contain thousands of health records and predict the risk of AD individually [14].

To address these issues, the field of large-scale and high-dimensional analytical pictures is receiving a lot of

interest in the developing field of deep learning, which customs underdone neuroscience facts to build functionality over "on-the-fly" training. We have reviewed publications systemically, using deep learning methods and neuroimaging information to detect AD early and predict AD progression [15]. ith this detailed overview, the intention of this learning is to provide readers with deep learning customized for AD and to help researchers to facilitate medical imaging. This work makes the main contribution

1. Introduced by an AI groundwork, AD is affected.
2. A clear road map for the need for deep learning and an analysis of general deep learning models for AD prediction has been provided in AD prediction.
3. Described the related research directions for medical researchers to make deep learning powered AD prediction.

This survey varies clearly from other recent surveys in the above points. It gives as much info as before. Section II discusses the theoretical source of AD to deep learning in AD. The paper has the following structure. Section III deals with a general data-driven deep learning model in AD prediction. Section IV lists and describes briefly the related directions of the research in Section V.

III. THEORETICAL BACKGROUND

3.1 Artificial Neural Networks, Transfer & Multi-kernel Learning

ANN is used extensively for modeling highly nonlinear data patterns in machine learning models. Conventional learning models use only one domain sample that greatly influences their performance when there are very few samples accessible. Transfer learning is a technique that employs samples not just from the target field, but also from other (related) subdomains. A multimodal multiple regularized transfer learning to help learn the target domain by transferring knowledge from a secondary area. Information from a nearby hospital is transmitted into samples using the TrAda Boost approach. Even when unconnected domains are disregarded, transferring knowledge from all source domains diminishes the model's performance. It's also possible that the sample labels supplied to them are inaccurate [1].

A robust approach has transformed original markings into multi-bit vectors and at the same time has gained common characteristics to identify unconnected domains. To tackle these challenges, The acquisition of data from specimens to local specimens through the subspace method is designed to solve these problems. EEG signals for different artificial neural network variants were used. Numerous variants of the artificial neural network, like the radial basis neural function network and the probabilistic neural network, have been studied and proven to be the best solution for neural vector quantization. The classification approach is developed among several various algorithms including Radial Based Neural Function Network, Best-First Decision Tree, Decision Tree, Labeling, and the Perceptron Multi layer.

A real-life event can't possibly contain all of the data needed to train a model. In these cases, the training system is insufficient. For resource classification, a self-adaptive network assignment classifier is employed. Primary component analysis is used for feature extraction and classification in order to deal with complex data [8]. Longitudinal information from MRI images is combined with a multi-layer perceptron and a two-directional gated recurrent unit. Information from synchronous auditory input is used by the long-term memory network for Alzheimer's disease categorization. With diffusion tensor pictures, a fluffy technique with the Artificial Neural Network is utilised to differentiate AD issues. Data from many modalities adds to the amount of information available to learning algorithms in machines [16].

For that purpose, a multi-kernel learning environment integrated into a common feature space by region of interest and tensor. In order to support classification, a new Laplacian multimodal, regularized smaller squares model is used to support unlisted samples. The selection features separately from each model ignore the strong intermodal correlation between individual subjects that lead to under-optimal performance. A new multi-task learning selection model was developed to resolve this difficulty, which combined sparse features from all modalities were selected. Until now, the various kernel learning methods listed linearly mix many kernels from diverse modalities and are particularly sensitive to each modality's weights. The non-linear combination of data from many modalities is used to solve these difficulties. Using a privileged evidence method, the training is utilised during the training phase. To make use of privileged information, a new restricted Boltzmann machine was proposed. For categorization, an ensemble method was used [5].

3.2 Pathophysiology of Alzheimer's Disease

Aside from the prefrontal, posterior, and parietal lobes, other brain regions contribute to our cognitive abilities, including the serotonin ventral nerve, neurotransmission cytosol, and choline cytosol. The hippocampal, amygdala, neocortex, and cerebral interconnections can all show signs of basal nuclei, neuronal atrophy, and/or illness. The pathogenesis of Alzheimer's illness is depicted in Figure 2.

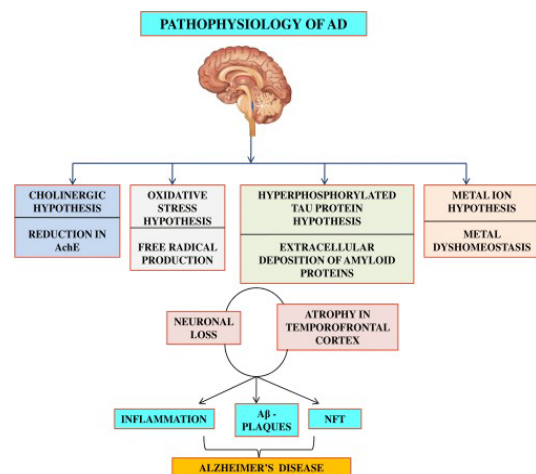


Fig. 2. Pathophysiology of Alzheimer's Disease

The hippocampal, the cerebral associating regions, the transentorhinal lobe, and the mesencephalon, all of which are influenced by the frontal, parietal, and occipital lobes, tend to develop tangles in a predictable order. The size and location of tangle formation is significantly more strongly associated with the ruthlessness of dementia than the magnitude of the development of senile plaques. Tau protein buildup has been associated to cognitive deterioration and cerebral shrinkage, particularly degeneration of the hippocampus. In Alzheimer's disease neuropathology, the temporofrontal cortex has neurons and atrophy loss, causing tenderness and depositing of amyloid plaques and an irregular cluster of portions of proteins, as well as a bundle of tangled fibres, resulting in an upsurge in the amount of cerebral cortex monocytes and macrophages, as well as stimulating the parenchymatous microglial cells in the parenchyma, resulting in an increase in the number of cerebral cortex monocytes [4].

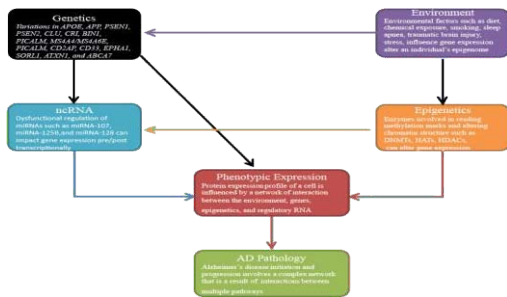


Fig. 3. Genetics, epigenetics and environmental influence - Alzheimer's disease pathology.

Ecological factors include peroxidation, inflammatory, and chemical contaminants. An individual's way of life can be affected by a variety of circumstances, including but not limited to their dietary habits and nutrition, brain trauma, inability to exercising, and smoke. AD pathology can be affected by genetic malfunction, non-coding of RNA and modifications in an individual's epigenome.[17].

IV. MACHINE LEARNING AND DEEP LEARNING MODELS FOR AD PREDICTION

Furthermore, inter-correlation-aware machine learning (ML) strategies have emerged as a pivotal component of computer-assisted statistical models, and are widely used in the computerized diagnostics and study of neurodevelopmental disorders. Although many different machine-learning approaches have been utilized for automated neurophysiological prediction and diagnosis, vector machine (SVM) and deep learning (DL)-based diagnostics approaches are two important avenues for further study. Numerous papers have been written about the application of machine learning techniques to diagnostic imaging. To highlight brain region correlations, functional connectivity (FC) patterns are widely used in contemporary SVM-based diagnostic models. Personalized FC arrangements are developed for paired regions of the brain that have been segregated using standard anatomical characteristics. In addition to its underperformance on original information, SVM has been criticised for the difficulty of obtaining significant features referring to [21].

As opposed to traditional models, DL platforms allow a computer to autonomously evaluate effective feature qualities in a training database using only the raw information supplied as input. The foundation of DL is an end-to-end modeled learning approach. End-to-end knowledge provides the advantage of simultaneously optimising all components of the optimization strategy, potentially resulting in maximum performance. In the realm of Alzheimer's disease prediction; Figure 4 depicts a general end-to-end hierarchy system model. There are four levels in the hierarchy, ranging from one (none) to four (everything) (full). The vast majority of current research uses Level 1 or Level 2 methods to achieve their objectives, which rely significantly on sophisticated software and, in some cases, precise parameter tweaks and manual noise removal. Because of these inter dependencies, those educators only used a part of the raw datasets for performance evaluation, leaving out obvious outliers and making it difficult to make a true enactment comparison. Additional benefit of end-to-end learning is that it allows you to observe how CNN arrived at their classification decision. The explanation aids in the clinician's understanding of CNN activity and the discovery of novel biomarkers. Level 2 explanations are restricted to the segmented area, which may obstruct understanding[22].

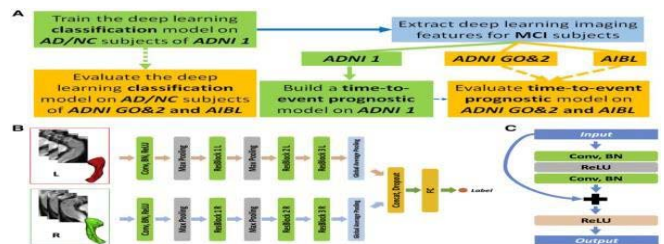


Fig. 4. Illustration of a generic, hierarchical system applicable to the field of Alzheimer's clinical diagnosis

Rather than building separate supervised models to forecast each attribute, a single model that predicts the evolution of numerous qualities can be built. One path to developing tools that can accurately simulate patient progression is to use statistical models based on artificial neural networks. Clinical data poses a number of challenges that are difficult to overcome with current machine learning algorithms. Many clinical datasets, for example, have a variety of data types (i.e., they are "multimodal"), few testing, and a high number of missed discoveries. When dealing with these issues, rigorous preprocessing or the deletion of variables that are too difficult to represent are frequently required. For example, one recent study focused solely on four characteristics that were frequently examined across 100,000 patients in a critical care unit data set on electronic health [13].

Three-dimensional (3D) or four-dimensional (4D) volume data, for example, is reformatted into 1D vector form and fed into DL networks such the limited Boltzmann machine (RBM) and deep belief network (DBN). Data complexity and time complexity, both of which are prominent components of medical data, can be blamed for the reliance on hand-crafted features. The ADNI dataset, for example, has only a few hundred images, yet each one has

over 10 million dimensions. It's important to remember that past methods misunderstood neighbourhood relationships (spatial proximity) in central nervous system information throughout the feature extraction phase. If spatial connections are not preserved, it is impossible to grasp a reliable description of how the network makes a categorization choice [20][9].

The convolutional neural network (CNN) has been proved to be a powerful DL model for grid-like data such as RGB and MR images. The use of CNNs has spread quickly across a wide range of disciplines, starting with AlexNet's amazing success on the natural picture categorization problem. Early advances in medical image processing were made with 2D pictures like retinal and chest X-rays, which were eventually expanded to 3D images like MRI. The majority of current CNN-based MRI methods are classified as Level 2. After segmenting the grey matter area for the period of pre-processing, several research employment it as a CNN input. Dropout, batch normalization, and residual module are three regularization strategies used in 3D-CNN-based approaches. While the use of effective regularization methods yielded promising results, no unsupervised learning was required. When faced with data shortages and increased dimensionality, unsupervised learning is considered important in the field of deep learning [19].

This represents the first efficient implementation of a volumetric CNN-based framework on MRI data employing 3D-stacked Convolutional Autoencoders for Alzheimer's disease classifications, but the prototype could only be emulated with an exactness of 80%. The models have been created using testing is to ensure data to anticipate temporal information and fine-tuning methodologies. Furthermore, there was little effort taken to justify the classifying choice. The use of Multiple Discourse Modalities in a Learning Environment techniques have tried to incorporate a variety of inputs and DL prototypes in order to address the optimization difficulty of AD. Grad-CAM uses different visual description approaches to demonstrate patchwise prediction inconsistencies in 3D-CNNs. Despite being capable to demonstrate how CNNs attained at their classification result, no endeavour has been through to address the progressive vs. stable MCI classification issue [3]. Shifting from variance. Given the complexity and multivariate nature of Alzheimer's disease, it's critical to simulate the evolution of whole patient profiles, including the progression of each sub-component of the ADAS-Cognitive and MMSE ratings, laboratory tests, and their correlations with baseline diagnostic criteria [9].

The development of strategies to overcome these restrictions is a critical step toward more widespread machine learning applications in precision medicine. Precision medicine is especially important for people with complex disorders who experience different disease progression and therapy responses. Alzheimer's illness and moderate cognitive injury (MCI) are both neurodegenerative diseases that cause a variety of cognitive and behavioral issues. To measure the severity of these symptoms, tests like the AD Valuation Gauge are frequently used. The heterogeneity of Alzheimer's illness and related dementia's makes them challenging to diagnose, control, and treat,

triggering calls for improved strategies for predicting and tracking disease development, as well as better clinical trial design for AD. Differential diagnosis is also interesting due to the difficulty of distinguishing between similar diseases.

V. COMPARISON ANALYSIS OF RELATED RESEARCH DIRECTIONS

Clinical evidence and imaging studies have been used to construct a number of disease progression models for MCI and AD. While previous methods for forecasting disease development were effective, they only predicted a specific outcome, such as the ADAS Cognitive score

Title	Classifier	Dataset	ADAccuracy	MCIAccuracy
[11]	RNN	PET	91.2	78.9
[12]	3DCNN	MRI	91.09	76.9
Lu et al. [2018]	DNN	MRI, PET	84.6	82.93
[6]	CNN	ADNI	95.73	82.31
[5]	Lenet-5, AlexNet, ZFNet, and R-CNN	MRI	75-25 crossvalidation and 90-10 crossvalidation are 97.68% and 98.75%	76.54
[18]	DNN	Kaggle	95.32	83.9

Patients with the same condition may experience various symptoms, progress at different rates, and respond to treatment in different ways. The basic goal of precision medicine is to figure out how to predict and treat differences between patients. Computational models of illness development based on machine learning and deep learning are an appealing way for dealing with patient heterogeneity. These computational models may one day be used to guide medical decisions; however, present implementations are limited by both the availability of data and the algorithms' capacity to extract insights from it.

VI. CONCLUSION

Supervised learning, a state-of-the-art machine learning approach, has surpassed classical machine learning in the area of computer vision by revealing deeper layers in complex, resistant to high. Applying Machine Learning to Predictive Health Care and automatic categorization of Alzheimer's disease (AD) has recently garnered a lot of interest, thanks to considerable advancements in neuroimaging that have yielded large-scale multimodal neuroimaging data. Multifunctional neuroimaging datasets used in conjunction with deep learning techniques for Alzheimer's disease diagnostics categorisation shows steady gains in terms of accuracy. Early-stage deep learning investigation in Alzheimer's disease aims to increase efficiency by using more hybrid types of information, such as -omics info, and boost openness by developing methods that are easy to explain and take into account domain experts' knowledge of the disease's unique features and paradigms. In this study, we analyse and compare different machine learning and deep learning data-driven designs based on massive organisational health information for AD risk stratification with the intention of better selecting

REFERENCES

- [1] Al-Shoukry, Suhad, Rassem, Taha and Makbol, Nasrin, "Alzheimer's Diseases Detection by Using Deep Learning Algorithms: A Mini-Review," IEEE Access, pp. 1-1, 2020, 10.1109/ACCESS.2020.2989396.
- [2] Ebrahimi, Amir and Chiong, Raymond, "Deep Learning to Detect Alzheimer's Disease from Neuroimaging: A Systematic Literature Review," Computer Methods and Programs in Biomedicine, vol. 187, p. 105242, 2019, 10.1016/j.cmpb.2019.105242.
- [3] Feng, Chiyu, Elazab, Ahmed, Yang, Peng, Wang, Tianshu, Zhou, Feng, Hu, Huoyou, Xiao, Xiaohua and Lei, Baiying, "Deep Learning Framework for Alzheimer's Disease Diagnosis via 3D-CNN and FSBi-LSTM," IEEE Access, pp. 1-1, 2019, 10.1109/ACCESS.2019.2913847.
- [4] Islam, Jyoti and Zhang, Yanqing, "Early Diagnosis of Alzheimer's Disease: A Neuroimaging Study with Deep Learning Architectures," pp. 1962-19622, 2018, 10.1109/CVPRW.2018.00247.
- [5] Janghel, and Rekh. "Deep-Learning-Based Classification and Diagnosis of Alzheimer's Disease", 2020, 10.4018/978-1-7998-0414-7.ch076.
- [6] Jain, Rachna, Jain, Nikita, Aggarwal, Akshay and D, Jude, "Convolutional Neural Network based Alzheimer's Disease Classification from Magnetic Resonance Brain Images," Cognitive Systems Research, p. 57, 2019, 10.1016/j.cogsys.2018.12.015.
- [7] Jo, Taeho, Nho, Kwangsik and Saykin, Andrew, "Deep Learning in Alzheimer's Disease: Diagnostic Classification and Prognostic Prediction Using Neuroimaging Data," Frontiers in Aging Neuroscience, p. 11, 2019, 10.3389/fnagi.2019.00220.
- [8] Ju, Ronghui, Hu, Chenhui, Zhou, Pan and Li, Quanzheng, "Early Diagnosis of Alzheimer's Disease Based on Resting-State Brain Networks and Deep Learning," IEEE/ACM Transactions on Computational Biology and Bioinformatics, pp. 1-1, 2017, 10.1109/TCBB.2017.2776910.
- [9] Dhanabalan, S. S., Sitharthan, R., Madurakavi, K., Thirumurugan, A., Rajesh, M., Avaniathan, S. R., & Carrasco, M. F. (2022). Flexible compact system for wearable health monitoring applications. Computers and Electrical Engineering, 102, 108130.
- [10] Liu, Siqi, Liu, Sidong, Cai, Weidong, Pujol, Sonia, Kikinis, Ron, Feng, and David Dagan Feng, "Early Diagnosis of Alzheimer's Disease with Deep Learning," 2014 IEEE 11th International Symposium on Biomedical Imaging, ISBI, pp. 1015-1018, 2014, 10.1109/ISBI.2014.6868045.
- [11] M. Liu, D. Cheng, and W. Yan, "Alzheimer's Disease Neuroimaging Initiative," Classification of Alzheimer's disease by combination of convolutional and recurrent neural networks using FDG-PET images, Front. Neuroinform. 12:35, 2018a.
- [12] M. Liu, J. Zhang, E. Adeli, and D. Shen, "Landmark-based deep multi-instance learning for brain disease diagnosis," Med. Image Anal., vol. 43, pp. 157-168, 2018b, 10.1016/j.media.2017.10.005.
- [13] Pazhani, A. A. J., Gunasekaran, P., Shanmuganathan, V., Lim, S., Madasamy, K., Manoharan, R., & Verma, A. (2022). Peer-Peer Communication Using Novel Slice Handover Algorithm for 5G Wireless Networks. Journal of Sensor and Actuator Networks, 11(4), 82.
- [14] Nguyen, Hoang and N. Chu, "An Introduction to Deep Learning Research for Alzheimer's Disease", IEEE Consumer Electronics Magazine, pp. 1-1, 2020, 10.1109/MCE.2020.3048254.
- [15] Ramzan, Farheen and Ghani, Usman and Rehmat, Asim and Iqbal, Sajid and Saba, Tanzila and Rehman, Amjad and Mehmood, Zahid, "A Deep Learning Approach for Automated Diagnosis and Multi-Class Classification of Alzheimer's Disease Stages Using Resting-State fMRI and Residual Neural Networks," Journal of Medical Systems, vol. 44, 2019, 10.1007/s10916-019-1475-2.
- [16] Salehi, and Waleed, "Alzheimer's Disease Diagnosis using Deep Learning Techniques," vol. 9, pp. 874-880, 2020, 10.35940/ijeat.C5345.029320.
- [17] So, Jae-Hong and Madusanka, Nuwan and Choi, Heung-Kook and Choi, Boo-Kyeong and Park, Hyeon-Gyun, "Deep Learning for Alzheimer's Disease Classification Using Texture Features", Current Medical Imaging Formerly: Current Medical Imaging Reviews, vol. 15, 2019, 10.2174/1573405615666190404163233.
- [18] Subramoniam, Manu, R. Aparna, R. Anurenjan and G. Sreeni, "Deep learning based prediction of Alzheimer's disease from magnetic resonance images", 2021.
- [19] K. Ananthajothi, and M. Subramaniam, "Multi level incremental influence measure based classification of medical data for improved classification," Cluster Comput., vol. 22, pp. 15073-15080, 2019, <https://doi.org/10.1007/s10586-018-2498-z>.
- [20] K. Ashokkumar, S. Parthasarathy, S. Nandhini, and K. Ananthajothi, "Prediction of grape leaf through digital image using FRCNN", Measurement: Sensors, vol. 24, p. 100447, 2022, ISSN 2665-9174, <https://doi.org/10.1016/j.measen.2022.100447>.
- [21] G. Balanagireddy and K. Ananthajothi, T.R. Ganesh Babu, and V. Sudha, "Correlation and Analysis of Overlapping Leukocytes in Blood Cell Images Using Intracellular Markers and Colocalization Operation," In AI Innovation in Medical Imaging Diagnostics, edited by Anbarasan, Kalaivani, Hershey, PA: IGI Global, pp. 137-154, 2021. <https://doi.org/10.4018/978-1-7998-3092-4.ch008>.
- [22] K. Ananthajothi, and M. Subramaniam, "Efficient Classification of Medical data and Disease Prediction using Multi Attribute Disease Probability Measure", Applied Mathematics and Information Sciences, ISSN 1935-0090, E-ISSN 2325-0399, Vol. 13, no. 5, pp. 783-789, 2019, <https://doi.org/10.18576/amis/130511>.