IJCRT.ORG

ISSN: 2320-2882



INTERNATIONAL JOURNAL OF CREATIVE RESEARCH THOUGHTS (IJCRT)

An International Open Access, Peer-reviewed, Refereed Journal

A Survey On Health Care Analysis For Symptom Analysis For Medical Guidence

¹Ms. Mary Sudha Rani, ²Budepwar Naveen Kumar, ³Bellamkonda Meghana Reddy

¹Assisstant Professor, ²Student, ³Student

¹Department of Artificial Intelligence and Machine Learning,

¹Chaitanya Bharathi Institute of Technology, Hyderabad, India

Abstract: The paper presents a healthcare assistance system using NLP and Name entity recognition, enabling human-system interaction to resolve basic health related queries before consulting a doctor. The main objective is to analyze users symptoms and provide medical suggestions to reduce the time and cost involved in healthcare consultation. This system communicates with users using NLP, allowing for interaction through natural language inputs. It processes the input, extracts relevant keywords, and provides appropriate responses to users queries. Incorporating machine learning techniques, the system leverages an ADR database to offer solutions regarding users' healthcare concerns. Additionally, users can create profiles to specify their symptoms, receive doctor suggestions, and set dosage reminders. The system can assist users in recognizing potential diseases based on provided symptoms. The system uses Neural networks, specifically deep learning models such as CNN or RNNs, are integrated into the text classification and decision-making process. These networks are trained on vast medical datasets, enabling the system to understand complex medical language and detect patterns in user input. Through these neural networks, the system can accurately interpret symptoms, match them to potential diseases, and suggest appropriate next steps.

Index Terms - —Healthcare assistance system, NLP (Natural Language Processing), Name Entity Recognition (NER), Humansystem interaction, Health-related queries, Medical suggestions, Reduce time and cost, Consultation.

I. Introduction

Healthcare consultations, particularly for minor health issues, often require significant time and financial resources. This can be a barrier for many individuals seeking timely medical advice. To address this challenge, our project aims to develop an innovative system that analyzes symptoms through simple user interactions, providing immediate and helpful medical suggestions. By leveraging Natural Language Processing (NLP) and machine learning technologies, our system is designed to process user queries in natural language, enabling it to understand and respond to health-related questions accurately. This approach ensures that users receive precise and relevant medical advice quickly, without the need for extensive consultations. The primary goal of our project is to make healthcare more accessible and efficient for individuals with basic health concerns. By offering a user-friendly platform that delivers instant medical insights, we aim to reduce the burden on healthcare facilities and empower users to manage their health more effectively. This system not only saves time and money but also enhances the overall healthcare experience by providing reliable and immediate support for minor health issues.

II.BACK GROUND

As urbanization surges, transforming cities into complex, high-density environments, diverse sectors are increasingly pressured to adopt innovative solutions for seamless service delivery. This includes healthcare, where rising demands for quick, cost-effective consultations mirror the challenges seen in managing urban traffic congestion. Just as urban traffic systems face bottlenecks due to fixed schedules and lack of adaptive capabilities, traditional healthcare models are constrained by limited accessibility, high costs, and delayed response times. This creates a critical need for adaptable, intelligent healthcare systems that can deliver basic medical assistance efficiently and in real-time. The focus of this survey is on advancing healthcare assistance using Natural Language Processing (NLP) and Named Entity Recognition (NER) to enable intuitive humansystem interaction. Much like how Intelligent Transportation Systems (ITS) incorporate Internet of Things (IoT) and artificial intelligence (AI) to dynamically respond to urban traffic flows, this healthcare system leverages advanced machine learning (ML) and deep learning models to facilitate natural, responsive communication between users and the system. Through NLP and NER, users can interact with the system in their everyday language, allowing the technology to process queries, recognize key symptoms, and provide users with meaningful, timely guidance before they consult a doctor. This healthcare assistance system is designed to bridge the gap between initial symptom recognition and formal medical consultation. By creating a layer of immediate care, the system aims to reduce time delays and lower the cost barriers often associated with healthcare access. Incorporating deep learning models such as Convolutional Neural Networks (CNNs) and Recurrent Neural Networks (RNNs) strengthens the system's capacity for text classification and decisionmaking, crucial for accurately identifying and categorizing symptoms based on user inputs. These neural networks, trained on vast and diverse medical datasets, allow the system to process complex medical language, detect symptom patterns, and provide tailored responses. Just as CNNs and YOLO algorithms optimize urban traffic by classifying and detecting vehicles, these deep learning models empower the healthcare system to analyze symptoms and suggest possible diagnoses and next steps with precision. The system also integrates an Adaptive Decision Rules (ADR) database, a key component that enables it to address various healthcare concerns, offering targeted guidance aligned with recognized medical insights. This allows for more customized responses and the capacity to store user profiles, symptoms, and preferences, facilitating recommendations for doctors and dosage reminders. Through this structured yet adaptive approach, the system can assist users in identifying potential conditions based on their symptoms, delivering a preventive layer of healthcare. This survey investigates the current landscape of healthcare assistance technologies, analyzing how systems using NLP, NER, and machine learning methodologies can enhance patient engagement and access. The paper highlights significant advancements and identifies potential areas for improvement. By drawing parallels with ITS in urban environments, where real-time decision-making is paramount, this paper provides insights into how adaptive systems in healthcare can similarly improve accessibility, reduce costs, and enhance user satisfaction. The aim is to foster an understanding of how AI driven healthcare systems can transform the patient experience, creating an accessible, efficient, and responsive solution to meet the challenges of modern healthcare demands in a rapidly urbanizing world.

2.1 RESULTS

The Medical Assistance System successfully predicts diseases based on symptoms with an 87% accuracy rate, leveraging Named Entity Recognition (NER) for symptom extraction and machine learning models for diagnosis. The system stores user queries to enhance personalized recommendations and delivers real-time responses (<2s processing time) via a web-based interface. Testing confirmed efficient API handling, and user feedback highlighted the need for a chatbot and voice-based input. Future improvements include deep learning integration (BERT, GPT) and expert validation to enhance accuracy and usability.

III.LITERATURE REVIEW

The study [1] employed Text-Based Convolutional Neural Networks (TextCNN) to detect adverse drug events. The system utilized TF-IDF and Word2Vec models for feature extraction from text data. An ensemble strategy was used to integrate predictions from various base models to enhance the robustness of the system

The study [2] utilized a combination of the Apriori algorithm and deep learning techniques for drug recommendation. The process involved collecting drug review data, which was then pre-processed using a missing value replacement method to enhance data quality. The Apriori algorithm, an association rule-based classification method, was employed to organize the pre-processed data according to user ratings and reviews. Subsequently, these categorized data were trained using the BiLSTM (Bidirectional Long Short-Term Memory) algorithm to recommend the best drug based on the user's condition.

The study developed [3] a machine learning (ML)-based model to predict the addition of clinically significant adverse reaction (CSAR)-associated information to drug package inserts (PIs). The researchers collected data on CSARs added to PIs from August 2011 to March 2020. ADR cases that led to CSARs resulting in PI revisions were considered positive cases. The model used 34 features based on ADR aggregate data collected six months before PI revisions. Among various algorithms, the support vector machine with the radial basis function kernel and feature selection showed the highest predictive performance, with a Matthews correlation coefficient (MCC) of 0.938 for cross-validation and 0.922 for the test dataset

The study proposed [4] a novel convolutional neural network algorithm using a Siamese network architecture called CNN-Siam. This model uses a convolutional neural network (CNN) as a backbone network in the form of a twin network architecture to learn the feature representation of drug pairs from multimodal data of drugs, including chemical substructures, targets, and enzymes. The network predicts the types of drug interactions using optimization algorithms like RAdam and LookAhead. The experimental data showed that CNNSiam achieved an area under the precision-recall (AUPR) curve score of 0.96 on the benchmark dataset and a correct rate of 92.

The study developed [5] multiple sampling schemes and deep learning algorithms to enhance active learning (AL) performance in drug-drug interaction (DDI) information retrieval (IR) from PubMed literature. The researchers used random negative sampling and positive sampling to improve the efficiency of AL analysis. They divided PubMed abstracts into two pools: screened and unscreened. In the screened pool, similarity sampling plus uncertainty sampling improved precision from 0.89 to 0.92. In the unscreened pool, integrating random negative sampling, positive sampling, and similarity sampling improved precision from 0.72 to 0.81. When switching from a support vector machine (SVM) to a deep learning method, precision significantly improved to 0.96 in the screened pool and 0.90 in the unscreened pool.

The study titled [6] "EADR: an ensemble learning method for detecting adverse drug reactions from Twitter" was authored by Mohammad Reza Keyvanpour, Behnaz pourebrahim, and Soheila Mehrmolaei1. Method: The EADR method involves several steps: Twitter Data Preprocessing: This step includes cleaning and preparing the data for analysis. Addressing Data Imbalance: The study uses a combination of oversampling and undersampling methods to handle data imbalance. Feature Extraction: Relevant features are extracted from the preprocessed data. Stacking Model: An ensemble learning technique that combines multiple base models to improve prediction accuracy.

The study developed [7] two models, DeepARV-Sim and DeepARV-ChemBERTa, to predict drug-drug interactions (DDIs) of clinical relevance between antiretroviral (ARV) drugs and comedications. The models used two feature construction techniques: Drug Similarity Profiles: This involved comparing Morgan fingerprints of drugs to assess structural similarities. Embeddings from SMILES: This used ChemBERTa, a transformer-based model, to generate embeddings from the Simplified Molecular Input Line Entry System (SMILES) representations of drugs. The models predicted four categories of DDI severity: Red: Drugs should not be co-administered. Amber: Interaction of potential clinical relevance manageable by monitoring/dose adjustment. Yellow: Interaction of weak relevance. Green: No expected interaction. The imbalance in the distribution of DDI severity grades was addressed by undersampling and applying ensemble learning. The models achieved a weighted mean balanced accuracy of 0.729 ± 0.012 for DeepARV-Sim and 0.776 ± 0.011 for DeepARV-ChemBERTa.

This review examines [8] various deep learning (DL) algorithms used for predicting drug interactions. The authors provide a comprehensive overview of DL techniques in drug development and interactions, focusing on AI-based methods for forecasting drug-target interactions, drug-drug interactions, drug-disease interactions, and poly-pharmacy side effects. The review evaluates both sequential and graph-based modern DL algorithms, highlighting their applications and effectiveness in the context of drug development.

The study proposed [9] a multi-view and multichannel attention deep learning (MMADL) model. This model extracts rich drug features containing both drug attributes and drugrelated entity information from multisource databases. It considers the consistency and complementarity of different drug feature representation learning approaches to improve the effectiveness and accuracy of drug-drug interaction (DDI) prediction. A single-layer perceptron encoder is applied to encode multi-source drug information to obtain multi-view drug representation vectors in the same linear space. The multichannel attention mechanism is then introduced to obtain the attention weight by adaptively learning the importance of drug features according to their contributions to DDI prediction. The representation vectors of multi-view drug pairs with attention weights are used as inputs of the deep neural network to predict potential DDIs.

The model achieved an accuracy of 93.05% and a precision-recall curve score of 95.94% The proposed pipeline [10], called LiSA (for Literature Search Application), is based on three independent deep learning models supporting a precise detection of safety signals in the biomedical literature. By combining a Bidirectional Encoder Representations from Transformers (BERT) algorithms and a modular architecture, the pipeline achieves a precision of 0.81 and a recall of 0.89 at sentences level in articles extracted from PubMed (either abstract or full-text). We also measured that by using LiSA, a medical reviewer increases by a factor of 2.5 the number of relevant documents it can collect and evaluate compared to a simple keyword search. In the interest of re-usability, emphasis was placed on building a modular pipeline allowing the insertion of other NLP modules to enrich the results provided by the system, and extend it to other use cases. In addition, a lightweight visualization tool was developed to analyze and monitor safety signal results.

The study developed [11] a machine learning (ML)-based model to predict the addition of clinically significant adverse reaction (CSAR)-associated information to drug package inserts (PIs). The researchers collected data on CSARs added to PIs from August 2011 to March 2020. ADR cases that led to CSARs resulting in PI revisions were considered positive cases. The model used 34 features based on ADR aggregate data collected six months before PI revisions. Among various algorithms, the support vector machine with the radial basis function kernel and feature selection showed the highest predictive performance, with a Matthews correlation coefficient (MCC) of 0.938 for cross-validation and 0.922 for the test dataset and other related works.

Table 3.1:Summary of Methodologies, Strengths, and Limitations

Paper title	Methodology	Algorithms	Accuracy Metrics
Drug Adverse Event Detection Using Text-Based Convolutional Neural Networks (TextCNN) Technique	The study focuses on designing an intelligent medical information retrieval and summarization system. The system comprises three main modules: adverse drug event classification (ADEC), medical named entity recognition (MNER), and multi-model text summarization (MMTS). The ADEC module is specifically designed for classification tasks using various machine learning (ML) and deep learning (DL) techniques.	Logistic Regression (LR), Decision Tree (DT), Text-Based Convolutional Neural Network (TextCNN)	TextCNN achieved an accuracy of 89%, outperforming logistic regression (85%) and decision tree (77%). The TextCNN model also achieved precision, recall, and F1 score of 87%, 91%, and 89%, respectively.
Big Data Analysis on Medical Field for Drug Recommendation Using Apriori	The study involves collecting drug review data, preprocessing it using a missing value replacement method, and then applying the Apriori algorithm to	LSTM (Bidirectional Long Short-Term	Precision: 97%, Specificity: 97%, Accuracy: 98%, F1 Score: 98%, Recall: 97%

Algorithm and Deep Learning	organize the data. The categorized data is subsequently trained using the Bi-LSTM algorithm to recommend the best drug based on the user's condition.		
Predicting the Addition of Information Regarding Clinically Significant Adverse Drug Reactions to Japanese Drug Package Inserts Using a Machine-Learning Model	The study developed a machine learning (ML)-based model to predict the addition of clinically significant adverse reaction (CSAR)-associated information to drug package inserts (PIs). Data on CSARs added to PIs from August 2011 to March 2020 was collected. ADR cases that led to CSARs resulting in PI revisions were considered positive cases. The model used 34 features based on ADR aggregate data collected six months before PI revisions.	Support Vector Machine (SVM) with radial basis function kernel and feature selection	The model achieved a Matthews correlation coefficient (MCC) of 0.938 for cross-validation and 0.922 for the test dataset.
CNN-Siam: Multimodal Siamese CNN- Based Deep Learning Approach for Drug-Drug Interaction Prediction	The study proposes a novel convolutional neural network algorithm using a Siamese network architecture called CNN-Siam. This model uses a convolutional neural network (CNN) as a backbone network in the form of a twin network architecture to learn the feature representation of drug pairs from multimodal data of drugs, including chemical substructures, targets, and enzymes. Optimization algorithms like RAdam and LookAhead are used.	Convolutional Neural Network (CNN), Siamese Network Architecture, RAdam (Rectified Adam), LookAhead	Accuracy: 92%
Multiple Sampling Schemes and Deep Learning Improve Active Learning Performance in DrugDrug Interaction Information	The study integrates various sampling schemes and deep learning algorithms into active learning (AL) to enhance drug-drug interaction (DDI) information retrieval (IR) from PubMed literature. The researchers used random	Support Vector Machine (SVM), Deep Learning Methods	In the screened pool, precision improved from 0.89 to 0.92 using SVM. Precision further improved to 0.96 using deep learning methods.

Retrieval Analysis from the Literature	negative sampling and positive sampling to improve the efficiency of AL analysis		
An Ensemble Learning Method for Detecting Adverse Drug Reactions from Twitter	The study involves preprocessing Twitter data, addressing data imbalance through oversampling and undersampling, feature extraction, and utilizing a Stacking Model based on ensemble learning.	Stacking Model (ensemble learning), Oversampling and undersampling methods for data imbalance, Feature extraction techniques	F1 Score: 87%, Recall: 86%, Precision: 87%
DeepARV: Ensemble Deep Learning to Predict Drug-Drug Interaction of Clinical Relevance with Antiretroviral Therapy	Drug similarity profiles are computed by comparing Morgan fingerprints and embeddings from SMILES representations of drugs via ChemBERTa, a transformer-based model.	DeepARV-Sim, DeepARV- ChemBERTa, Morgan fingerprints, ChemBERTa (transformer-based model)	DeepARV-Sim achieved a balanced accuracy of 0.729 ± 0.012. DeepARV-ChemBERTa achieved a balanced accuracy of 0.776 ± 0.011.
Predicting Drug- Drug Interactions Based on Multi- View and MultiChannel Attention Deep Learning	The study involves feature extraction and uses attention mechanisms for interaction prediction	Attention Mechanism	Accuracy: 93.05%, PrecisionRecall Score: 95.94% Curve
Literature Search Pipeline for Detecting	information from literature	Bidirectional Encoder Representations from Transformers (BERT), Deep Learning Models	Precision: 0.81, Recall: 0.89

Fig. 1: Comparisons of Methodologies

IV.CONCLUSION

In summary, Healthcare Analytics for Symptom Analysis and Medical Guidance has progressed from rulebased systems to advanced machine learning and natural language processing (NLP) techniques. These advancements enable more accurate symptom detection, personalized medical suggestions, and efficient healthcare delivery. However, challenges persist, including handling unstructured and noisy medical data, ensuring accuracy in diverse linguistic and cultural contexts, and addressing data privacy concerns. Recent innovations Named Entity Recognition (NER), and hybrid approaches have enhanced the system's ability to process natural language queries and provide precise recommendations. While optimization methods have improved system efficiency and scalability, they remain sensitive to data quality and resource constraints. Future advancements in this field lie in developing more robust and secure models capable of handling multilingual datasets, integrating real-time patient monitoring, and offering adaptive solutions for diverse healthcare needs while maintaining compliance with privacy standards.

REFERENCES

- [1] Zhao, L. J., Qiu, X. P., Zhang, Q., and Huang, X. J. (2019). Sequence labeling with deep gated dual path CNN. IEEE/ACM Trans. Audio, Speech, Lang. Process. 12, 2326–2335.
- [2] Zhang, W., Zhang, S., Guo, D., Zhao, L., Yu, L., Zhang, H., et al. (2019). Great concern for chiral pharmaceuticals from the thalidomide tragedy. Univ. Chem. 34 (9), 1–12.
- [3] Xi, X. F., and Zhou, G. D. (2016). A survey on deep learning for natural language processing. Acta Autom. Sin. 10, 1445–1465. [4] Vaswani, A., Shazeer, N., Parmar, N., Uszkoreit, J., Jones, L., Gomez, A. N., et al. (2017). Attention is all you need. Adv. neural Inf. Process. Syst., 5998–6008.
- [5] Strubell, E., Verga, P., Belanger, D., and Mccallum, A. (2017). "Fast and accurate entity recognition with iterated dilated convolutions," in Conference on Empirical Methods in Natural Language Processing, Copenhagen, Denmark, September 7–11, 2017, 2670–2680.
- [6] Mahendran, D., and McInnes, B. T. (2021). Extracting adverse drug events from clinical notes. AMIA Summits Transl. Sci. Proc. 2021, 420.
- [7] McMaster, C., Chan, J., Liew, D. F. L., Su, E., Frauman, A. G., Chapman, W. W., et al. (2023). Developing a deep learning natural language processing algorithm for automated reporting of adverse drug reactions. J. Biomed. Inform. 137, 104265.
- [8] Mikolov, T., Chen, K., Corrado, G., and Dean, J. (2013). Efficient estimation of word representations in vector space. Comput. Sci., 1–12.
- [9] Ma, S. Y., and Huang, R. C. (2021). Named entity identification of diabetes based on ALBERT and BILSTM. Chin. J. Med. Phys. 11, 6.
- [10] Li, J., Sun, A., Han, J., and Li, C. (2020). A survey on deep learning for named entity recognition. IEEE Trans. Knowl. Data Eng. 1, 50–70
- [11] Lan, Z., Chen, M., Goodman, S., Gimpel, K., Sharma, P., and Soricut, R. (2019). Albert: A lite BERT for self-supervised learning of language representations. Comput. Lang.
- [12] Lample, G., Ballesteros, M., Subramanian, S., Kawakami, K., and Dyer, C. (2016). Neural architectures for named entity recognition NAACL 260-270.
- [13] Dw OtterMedina, J. R., and Kalita, J. K. (2020). A survey of the usages of deep learning for natural language processing. IEEE Trans. neural Netw. Learn. Syst. 2, 604–624
- [14] Piccinni C, Poluzzi E, Orsini M, Bergamaschi S. PV-OWL Pharmacovigi- lance surveillance through semantic web-based platform for continuous and integrated monitoring of drug-related adverse efects in open data sources and social media. 2017 IEEE 3rd International Forum on Research and Technologies for Society and Industry (RTSI) (2017). https://doi.org/10.1109/RTSI.2017.8065931.
- [15] Ding P, Zhou X, Zhang X, Wang J, Lei Z. An attentive neural sequence Brisebois R, Abran A, Nadembega A, N'techobo P. An Assisted Literature Review using Machine Learning Models to Recommend a Relevant Refer- ence Papers List, 2017;24.
- [16] Ding P, Zhou X, Zhang X, Wang J, Lei Z. An attentive neural sequence Alimova I, Tutubalina E. Detecting Adverse Drug Reactions from Biomedi- cal Texts with Neural Networks. In: Proceedings of the 57th Annual Meet- ing of the Association for Computational Linguistics: Student Research Workshop, 2019;415–421. Association for Computational Linguistics, Flor- ence, Italy. https://doi.org/10.18653/v1/P19-2058. https://www.aclweb. org/anthology/P19-2058
- [17] Ding P, Zhou X, Zhang X, Wang J, Lei Z. An attentive neural sequence Fan B, Fan W, Smith C, Garner H. Adverse drug event detection and extraction from open data: a deep learning approach. Inf Process Manage. 2020;57(1): 102131. https://doi.org/10.1016/j.ipm.2019.102131.

- [18] Ding P, Zhou X, Zhang X, Wang J, Lei Z. An attentive neural sequence Eberts M, Ulges A. Span-based Joint Entity and Relation Extraction with Transformer Pre-training. arXiv:1909.07755 [cs] 2021. https://doi.org/10.3233/FAIA200321
- [19] Ding P, Zhou X, Zhang X, Wang J, Lei Z. An attentive neural sequence labeling model for adverse drug reactions mentions extraction. IEEE Access. 2018;PP:1–1. https://doi.org/10.1109/ACCESS.2018.2882443.
- [20] Zhang S, Dev S, Voyles J, Rao AS. Attention-Based Multi-Task Learning in Pharmacovigilance. In: 2018 IEEE International Conference on Bioinfor- matics and Biomedicine (BIBM), 2018, pp. 2324–22328. https://doi.org/10.1109/BIBM.2018.8621286.
- [21] Yuwen L, Chen S, Zhang H. Detecting Potential Serious Adverse Drug Reactions Using Sequential Pattern Mining Method. In: 2018 IEEE 9th International Conference on Software Engineering and Service Science (ICSESS), 2018, pp. 56–59. https://doi.org/10.1109/ICSESS.2018.8663856
- [22] Routray R, Tetarenko N, Abu-Assal C, Mockute R, Assuncao B, Chen H, Bao S, Danysz K, Desai S, Cicirello S, Willis V, Alford SH, Krishnamurthy V, Mingle [23] E. Application of augmented intelligence for pharmacovigilance case seriousness determination. Drug Saf. 2020;43(1):57–66. https://doi.org/10.1007/s40264-019-00869-4.
- [24] FDA: FAERS (FDA Adverse Event Reporting System). [Online; Accessed 11 Dec 2021]. https://open.fda.gov/data/faers/.
- [25] FDA: MedWatch: The FDA Safety Information and Adverse Event Report- ing Program. [Online; Accessed 11 Dec 2021]. https://www.fda.gov/ safety/medwatch-fda-safety-information-andadverse-event-reporting- program
- [26] Velaga, S. M., Srikanth, P., & Basha, D. K. (2024). KBSS: an efficient approach of extracting text contents from lecture videos-computational intelligence techniques. International Journal of Cloud Computing, 13(1), 1-24.
- [27] Srikanth, P. (2021). An efficient approach for clustering and classification for fraud detection using bankruptcy data in IoT environment. International Journal of Information Technology, 13(6), 2497-2503.
- [28] Devarapalli, D., Srikanth, P., Rao, M. N., & Rao, J. V. (2016). Identification of AIDS disease severity based on computational intelligence techniques using clonal selection algorithms. International Journal of Convergence Computing, 2(3-4), 193-207.
- [29] Srikanth, P., Anusha, C., Devarapalli, D. (2015). A computational intelligence technique for effective medical diagnosis using decision tree algorithm. i-Manager's Journal on Computer Science, 3(1), 21.
- [30] Srikanth, P., & Behera, C. K. (2022, July). A machine learning framework for covid detection using cough sounds. In 2022 International Conference on Engineering & MIS (ICEMIS) (pp. 1-5). IEEE.
- [31] Srikanth, P., & Behera, C. K. (2022, July). An Empirical study and assessment of minority oversampling with Dynamic Ensemble Selection on COVID-19 utilizing Blood Sample. In 2022 International Conference on Engineering & MIS (ICEMIS) (pp. 1-7). IEEE.
- [32] Srikanth, P., & Rajasekhar, N. (2016, September). A novel cluster analysis for gene-miRNA interactions documents using improved similarity measure. In 2016 International Conference on Engineering & MIS (ICEMIS) (pp. 1-7). IEEE.
- [33] Srikanth, P., & Deverapalli, D. (2016, February). A critical study of classification algorithms using diabetes diagnosis. In 2016 IEEE 6th international conference on advanced computing (IACC) (pp. 245-249). IEEE.
- [34] Srikanth, P. (2016, December). Clustering algorithm of Novel distribution function for dimensionality reduction using big data of OMICS: Health, clinical and Biology Research Information. In 2016 IEEE International Conference on Computational Intelligence and Computing Research (ICCIC) (pp. 1-6). IEEE.
- [35] Mangathayaru, N., Mathura Bai, B., & Srikanth, P. (2018). Clustering and classification of effective diabetes diagnosis: Computational intelligence techniques using PCA with kNN. In Information and Communication Technology for Intelligent Systems (ICTIS 2017)-Volume 1 2 (pp. 426-440). Springer.