

WELCOME ALL !!



# BIOLX

A Biomedical Named Entity Recognition App

10<sup>th</sup> May 2024, Friday

# OUR GUIDE

Ms. Anupama M L

Asst Prof of CSE Department

## TEAM MEMBERS

Akhilesh M S

S8 CSE Student

Varun Sasikumar

S8 CSE Student

Rintu Sam

S8 CSE Student

Rishal Mohammed V A

S8 CSE Student

# TABLE OF CONTENTS

01

Introduction

02

Existing  
System

03

Objectives

04

Literature  
Survey

04

# TABLE OF CONTENTS

05

System  
Diagrams

06

Produced  
Output

07

Conclusion

08

References

# 01 Introduction

# WHAT IS BioNER ?

- ✓ BioNER is a subfield of natural language processing (NLP) that aims to identify and classify biomedical entities in text.
- ✓ Biomedical entities are specific types of words or phrases that refer to biomedical concepts, such as genes, proteins, diseases, drugs, and chemicals.

# WHY IS BioNER IMPORTANT ?

- ✓ BioNER is important because it enables the extraction of biomedical knowledge from unstructured text, which can be used to support a wide range of applications, such as drug discovery, clinical research, and public health surveillance.

# FOR EXAMPLE

Biologically Active Substance

Drug

- ✓ The benefits of taking cholesterol lowering statin drugs outweigh the risks even among people who are likely to develop diabetes.

Disorder

Organic chemical

Enzyme

- ✓ Acute exposure to resveratrol inhibits AMPK activity in human skeletal muscle cells.

Cell



02

# Existing System

# EXISTING SYSTEM

- ✓ Some existing systems employ rule-based approaches for entity recognition, which may lack flexibility in handling complex medical terminology and variations.
- ✓ The existing system may utilize traditional NLP techniques such as tokenization and part-of-speech tagging, which may not adequately capture the nuances of biomedical text.

# DISADVANTAGES

- ✓ Inaccuracy
- ✓ Scalability Challenges
- ✓ Expert Dependency
- ✓ Lack of Real-Time Processing
- ✓ Limited Adaptability

# PROBLEM DESCRIPTION

- ✓ Manual Effort
- ✓ Limited Handling
- ✓ Language Complexity

# PROPOSED SYSTEM

- ✓ The development of a mobile application for Biomedical Named Entity Recognition (BioNER) represents a significant advancement in the domain of biomedical research and healthcare.
- ✓ This specialized mobile app harnesses a state-of-the-art BioNER model hosted on a cloud server, offering a range of features designed to enhance efficiency, accuracy, understanding, and accessibility for biomedical professionals and researchers.

# 03 Objectives

# OBJECTIVES

- ✓ Develop a biomedical entity recognition system using CRF and BERT models.
- ✓ Create an intuitive Flutter mobile app for users to input biomedical text.
- ✓ Implement RESTful APIs with Flask to facilitate communication between the frontend and backend.
- ✓ Integrate CRF and BERT models to enhance entity recognition accuracy.
- ✓ Document the development process and create a reliable, user-friendly application.

# 04 Literature Survey

# PAPER 01

Name	Authors	Abstract	Proposed System
<b>Biomedical Named Entity Recognition via Reference-Set Augmented Bootstrapping</b>	<ul style="list-style-type: none"><li>✓ Joel Mathew</li><li>✓ Shobeir Fakhraei</li><li>✓ Jose Luis Ambite</li></ul>	<ul style="list-style-type: none"><li>✓ Train a neural NER model on a small seed of labeled data.</li><li>✓ Use a reference set of entity names to identify entity mentions with high precision but low recall on an unlabeled corpus.</li><li>✓ Use the NER model to assign weak labels to the corpus and retrain the model iteratively on the augmented training set.</li></ul>	<ul style="list-style-type: none"><li>✓ Create an augmented training set using a small labeled seed set and a large unlabeled corpus.</li><li>✓ Use automated labeling with a reference set to weakly label the unlabeled corpus.</li><li>✓ Apply an iterative label refinement process.</li></ul>

# PAPER 02

Name	Authors	Abstract	Proposed System
<b>Bio-NER: Biomedical Named Entity Recognition using Rule-Based and Statistical Learners</b>	<ul style="list-style-type: none"><li>✓ Pir Dino Soomro</li><li>✓ Sanotsh Kumar</li><li>✓ Banbhrani</li><li>✓ Arsalan Ali Shaikh</li><li>✓ Hans Raj</li></ul>	<ul style="list-style-type: none"><li>✓ The authors propose a hybrid approach for biomedical NER, employing rule-based and statistical methods.</li><li>✓ They introduce a novel method for disease NER that surpasses existing methods.</li><li>✓ The effectiveness of their approach is validated using standard evaluation metrics.</li></ul>	<ul style="list-style-type: none"><li>✓ The proposed hybrid approach for biomedical NER combines machine learning, rule-based, and decision tree methods.</li><li>✓ It uses simple features like affixes and context, as well as more complex features like orthographic, POS tags, and N-grams.</li></ul>

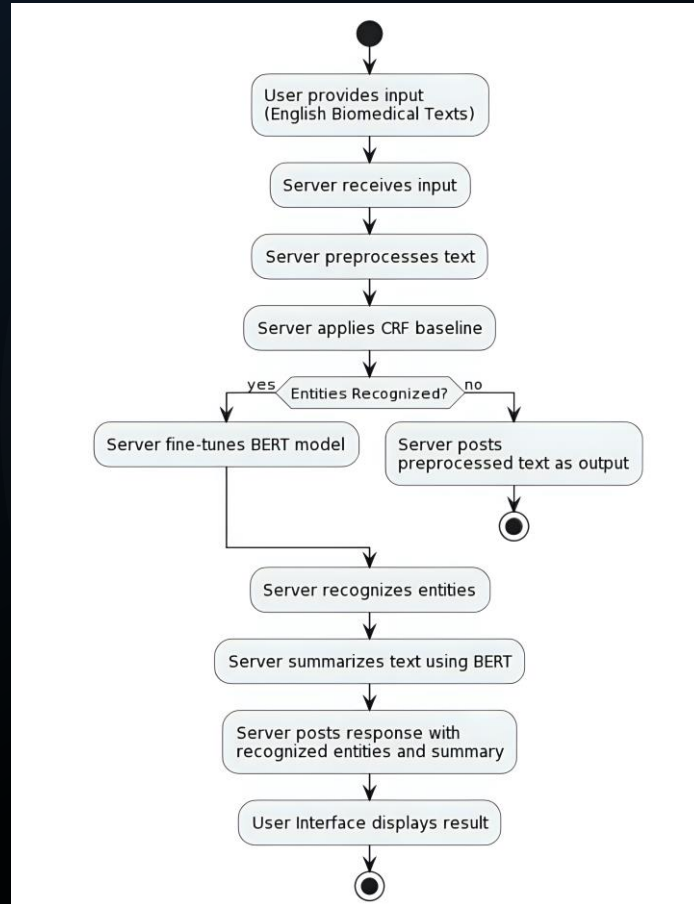


# PAPER 03

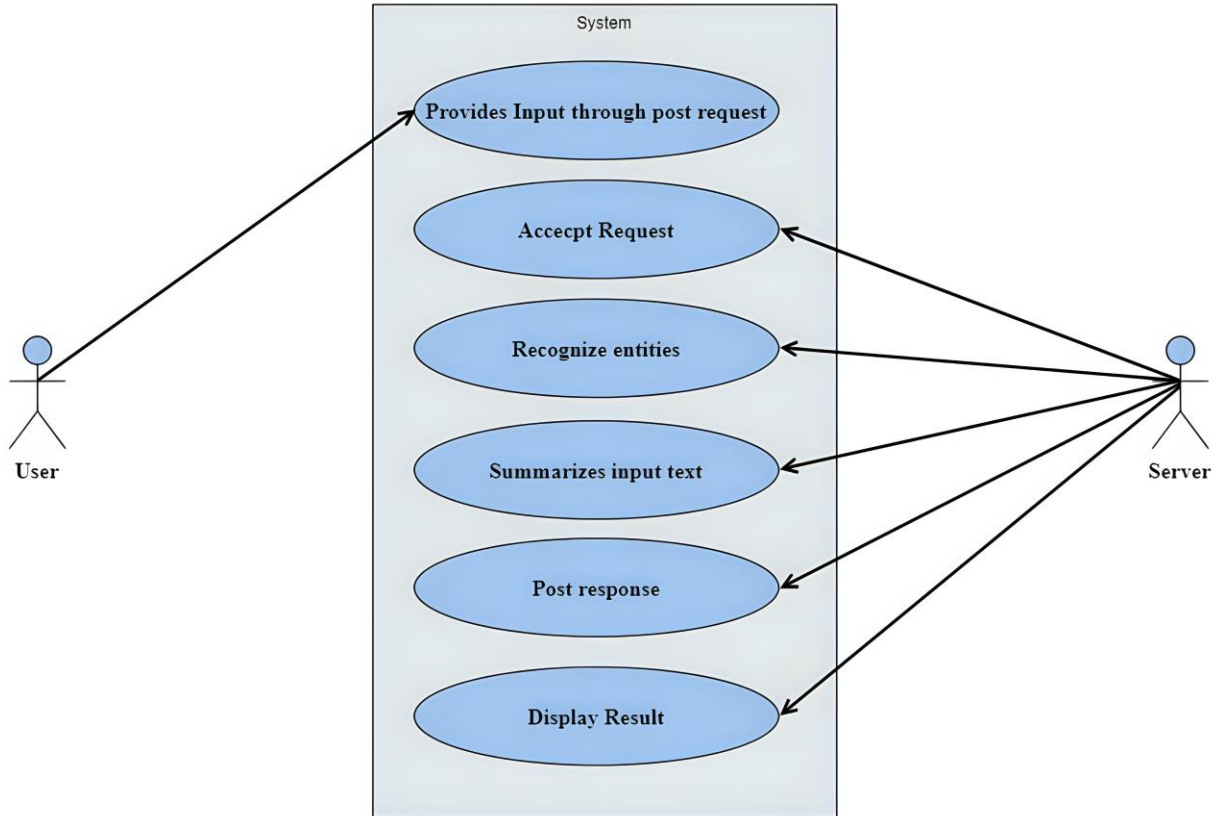
Name	Authors	Abstract	Proposed System
<b>Unsupervised biomedical named entity recognition: Experiments with clinical and biological texts</b>	<ul style="list-style-type: none"><li>✓ Shaodian Zhang</li><li>✓ Noémie Elhadad</li></ul>	<ul style="list-style-type: none"><li>✓ Stepwise approach to entity boundary detection and entity type classification.</li><li>✓ Uses noun phrase chunking and distributional semantics principles, without relying on handcrafted rules or annotated data.</li><li>✓ Outperforms a baseline dictionary match approach on clinical and biological texts.</li></ul>	<ul style="list-style-type: none"><li>✓ Extracts candidate entities from free text using a noun phrase chunker and a filter based on inverse document frequency.</li><li>✓ Classifies candidate entities into categories of interest using principles from distributional semantics.</li><li>✓ Does not rely on handcrafted rules or annotated data.</li></ul>

# 05 System Diagrams

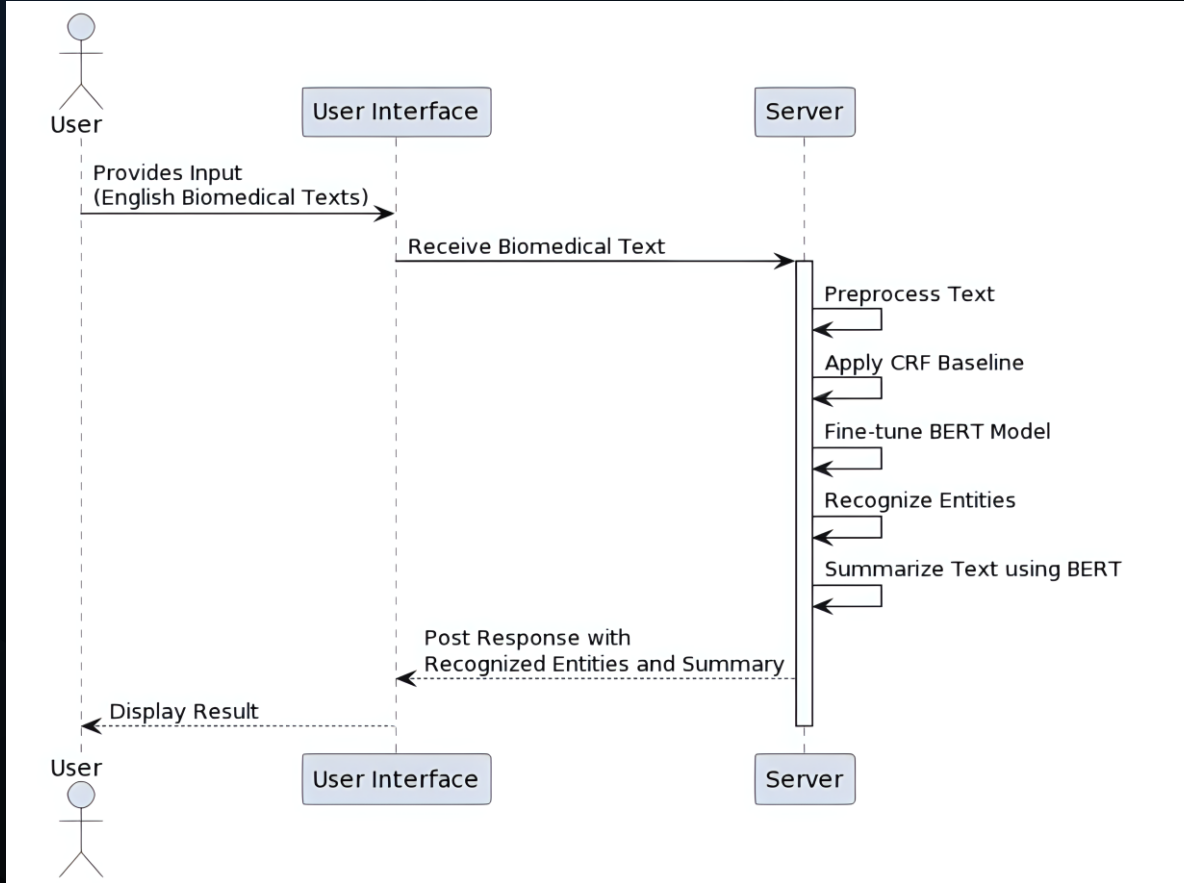
# ACTIVITY DIAGRAM



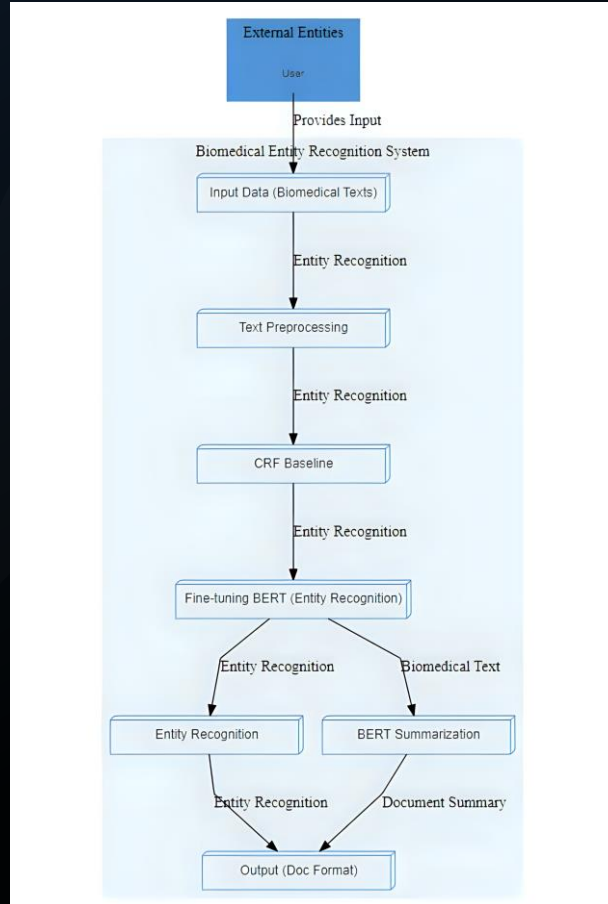
# USE CASE DIAGRAM



# SEQUENCE DIAGRAM



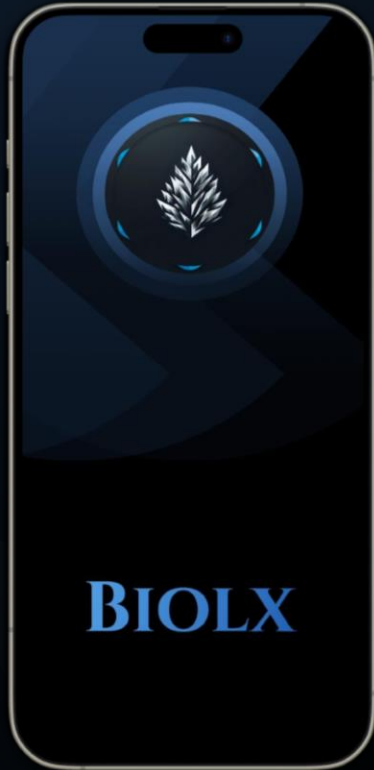
# DATAFLOW DIAGRAM



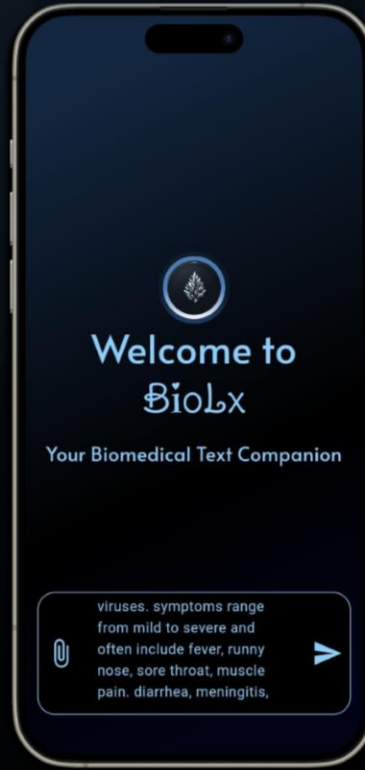
06

# Produced Output

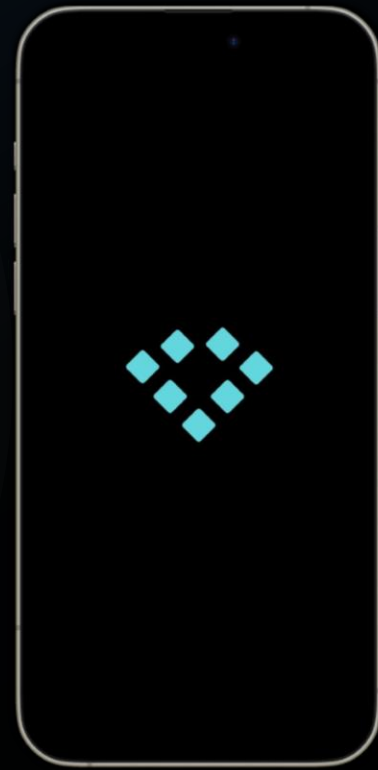
SPLASH SCREEN



HOME SCREEN

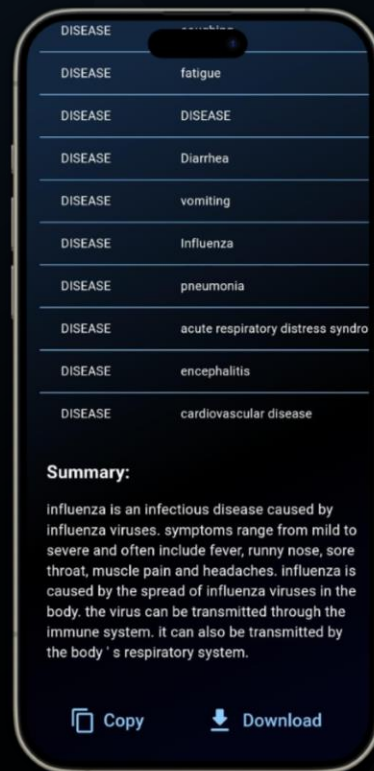


LOADING SCREEN





# DISPLAY SCREEN



# 07 Conclusion

# CONCLUSION

- ✓ The biomedical entity recognition project combines CRF and BERT models for accurate identification of entities in biomedical texts.
- ✓ Flask as the primary backend framework, and Python as the programming language supporting the overall backend functionality.
- ✓ The team work between CRF and BERT captures both local and global context, enhancing entity recognition.
- ✓ Avoiding a database simplifies the system, utilizing in-memory knowledge and alternative storage effectively.
- ✓ This system offers a practical, user-friendly solution for biomedical text processing, with potential for scalability and knowledge base refinement in future updates.

08

# References

# REFERENCES

- ✓ Gamal Crichton, Sampo Pyysalo, Billy Chiu, and Anna Korhonen. 2017. A neural network multi-task learning approach to biomedical named entity recognition. BMC bioinformatics, 18(1):368.
- ✓ Jacob Devlin, Ming-Wei Chang, Kenton Lee, and Kristina Toutanova. 2018. BERT: Pre-training of deep bidirectional transformers for language understanding. arXiv preprint arXiv:1810.04805
- ✓ Diederik P Kingma and Jimmy Ba. 2014. Adam: A method for stochastic optimization. arXiv preprint arXiv:1412.6980.

THANK YOU !!