

Al-Powered Leukemia Detection

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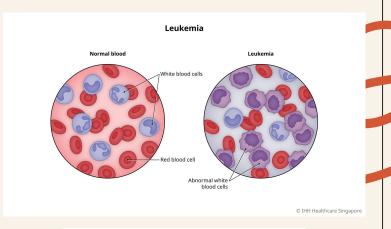


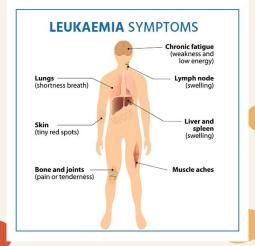
Project Overview

- Leukemia is a type of blood cancer that affects white blood cells, and early detection is critical for effective treatment.
- Our project focuses on using machine learning to improve the early diagnosis of leukemia by analyzing blood test results.

Problem Statement

- Leukemia symptoms including fatigue, bruising, and bone pain are vague and often misdiagnosed or detected too late, especially in chronic cases.
- Delayed diagnosis leads to more invasive treatments and lower survival rates.
- There is a critical need for faster, more accurate diagnostic tools so that patients can begin receiving treatment sooner.





Significance

- ~24,000 deaths from leukemia occur annually in the U.S.
- Blood plays a critical role in homeostasis and immune function.
- Cancer in the bloodstream can rapidly affect multiple systems.
- Earlier diagnosis allows for earlier treatment which leads to less invasive, more effective outcomes.
- Biomedical engineers aim to revolutionize diagnosis using machine learning & data analysis for faster, more accurate detection.



Data

- Dataset included various biomarkers: individual's lactate dehydrogenase levels, white blood cell count, gene expression, and the CRP protein levels.
- Data was labeled with either a 0 (false) or 1 (true) to denote whether or not the patient had leukemia.
- Dataset was synthetic (artificially generated), but based on real, accurate values of biomarker levels in leukemia and non-leukemia patients.

Methodology

Our code:



- Loads the synthetic dataset
- Explores the dataset



 Preprocesses the data, keeping all relevant features (all features were relevant in this case)



Trains the models and tunes settings for best precision



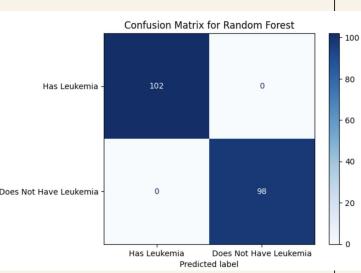
 Evaluates the model performance (confusion matrix, accuracy, precision, recall, and F1 score)



 Takes new user input and predicts leukemia status using trained model

Results

- Accuracy
 - Definition: percentage of correct predictions
 - o Our model: 100% accurate
- Precision
 - Definition: accuracy of model's positive predictions
 - Our model: 100% precise (no missed diagnosis)
- Recall
 - Definition: percent of true positives identified by the model
 - Our model: 100% recall (no false negatives)
 - F1 Score
 - Definition: harmonic mean of precision and recall
 - Our model: 100%



Link to Code

Github Repository Link