



# **Medline Search Engine (MSE)**

**CIS 612 – Big Data & Analytics**

**Final Project Report**

*Submitted By:*

Dinky Mishra – 2864923

Alim Khan Abdul – 2882808

Sushma Reddy Avala – 2885387

Aditya Sairam Pullabhatla – 2863159

## Table of Contents

- 1. Data Description, Data Size, and Data Collection Method**
  - Data Description
  - Data Size
  - Data Collection Method
  - Data Storage in MongoDB and MySQL
- 2. Goal of the Intelligent Big Data Analytic Application**
  - Project Goals
  - AI Components: Natural Language Processing and Machine Learning Algorithms
- 3. Platform Setting and System Configuration**
  - Development Environment
  - Node.js Environment Setup
  - MongoDB and MySQL Setup
  - Frontend Setup and Server Configuration
- 4. System Design and Architecture**
  - Overall System Architecture
  - Data Flow Diagram
- 5. Raw Big Data Preprocessing Methods**
  - Data Cleaning (cleanData.js)
  - Text Normalization
  - Intermediate Results and Sample Cleaned Data Snapshot
- 6. Big Data Processing Pipeline and Data Transformation**
  - Pipeline Steps: Data Ingestion, Tokenization, Normalization
  - Feature Extraction and Inverted Index Creation
  - Data Transformation Methods
  - Inverted Index and Dictionary Table Snapshots
- 7. Knowledge Base Structure and Database Design**
  - MongoDB Schema for Raw Data
  - MySQL Schema for Processed Data
  - Knowledge Base Features
- 8. Scoring and Ranking Algorithms**
  - TF-IDF (Term Frequency-Inverse Document Frequency)
  - Cosine Similarity
- 9. Problems Encountered and Resolutions**
  - XPath Issues in Web Scraping
  - Inefficient Database Query Execution
- 10. System Demo**
  - User Interface
  - Backend Search Logic and Execution Steps
  - Search Demo and Results Snapshots
- 11. Execution Steps**
  - Backend and Data Collection Setup
  - Frontend Setup for Search Engine UI
  - Code Overviews for Key Scripts
- 12. Conclusion**

# 1. Data Description, Data Size, Data Collection Method

## Data Description

The Medline Search Engine project aimed to create a comprehensive search platform for medical information by leveraging data from the MedlinePlus Encyclopedia. The dataset contains detailed medical articles, which include:

- **Article Titles:** Capturing the main topic of each article.
- **URLs:** Providing links to the original articles on the MedlinePlus platform.
- **Content:** Textual content covering medical topics, including diseases, symptoms, procedures, medications, and medical conditions. The content was extracted to facilitate efficient indexing and querying.

## Data Size

- **Total Articles Scraped:** 4,500 approx. articles from MedlinePlus.
- **Number of Terms:** Over 1.4 million terms after preprocessing, including specialized medical terms and general content terms.

## Data Collection Method

Data collection was accomplished through a custom web scraper developed using Node.js and Puppeteer. This scraper was designed to efficiently navigate and extract content from the MedlinePlus website.

- **Tool Used:** webScraper.js

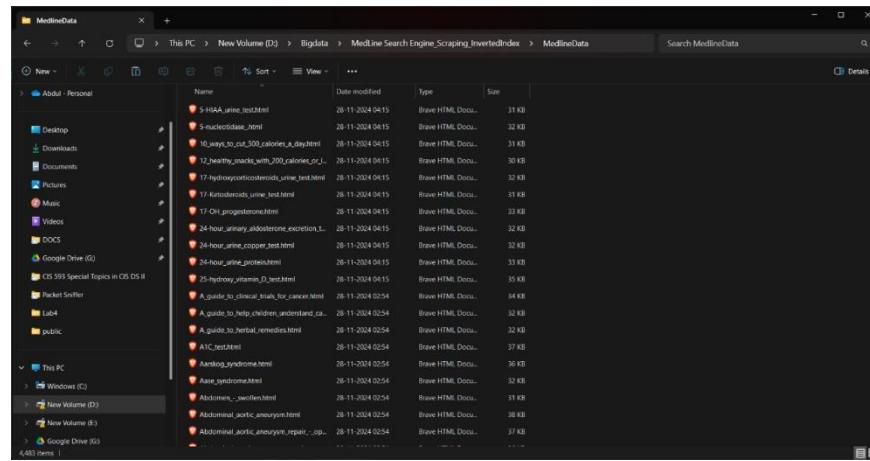
```
import puppeteer from "puppeteer";
import fs from "fs/promises";

const main = async () => {
  const browser = await puppeteer.launch();
  const page = await browser.newPage();
  await page.goto("https://medlineplus.gov/encyclopedia.html");
  // Scrape data logic here
  await browser.close();
};

await main();
```

- **Process:**

- **Automated Scraping:** The script navigated the MedlinePlus Encyclopedia website, accessed the articles, and extracted key details such as titles, URLs, and HTML content.
- **Data Storage:** Scrapped data was stored initially as raw HTML files and then imported into MongoDB for efficient retrieval and backup purposes.
- **Data Validation:** Data accuracy and completeness were validated by cross-referencing the number of scraped articles with the total number of articles available on the MedlinePlus website. Duplicate articles were removed, and inconsistencies were handled to maintain data integrity.



## Medline Data Snapshot in MongoDB:

- **Description:** This screenshot shows a preview of the data stored in the medlinedata table in MongoDB. It displays multiple records, including fields such as id, title, href, fileContent and filePath providing an overview of the unstructured data.

_id	title	href	fileContent	filePath
_id: ObjectId('674834ef52bed91ebedca05')	"A guide to clinical trials for cancer"	"https://medlineplus.gov/ency/patientinstructions/0008823.htm"	"<DOCVTP> html><html lang="en"> <div> <h2>A guide to clinical trials for cancer</h2> ...</div> <div> <p>This page is a guide to clinical trials for cancer. Clinical trials are studies that test new ways to prevent, detect, or treat cancer. ...</p> <p>Clinical trials are conducted at many medical centers across the country. ...</p> <p>If you're interested in participating in a clinical trial, you can contact your doctor or a medical center near you. ...</p>"	"D:\Bigdata\Medline Search Engine_Scraping_InvertedIndex\MedlineData\A_0008823.htm"
_id: ObjectId('674834ef52bed91ebedca05')	"A guide to herbal remedies"	"https://medlineplus.gov/ency/patientinstructions/0008808.htm"	"<DOCVTP> html><html lang="en"> <div> <h2>A guide to herbal remedies</h2> ...</div> <div> <p>Herbal remedies are substances made from plants or parts of plants. ...</p> <p>Some people use herbal remedies to treat health problems. ...</p> <p>It's important to talk to your doctor before taking any herbal remedy. ...</p>"	"D:\Bigdata\Medline Search Engine_Scraping_InvertedIndex\MedlineData\A_0008808.htm"
_id: ObjectId('674834ef52bed91ebedca05')	"AIC"	"https://medlineplus.gov/ency/article/0008849.htm"	"<DOCVTP> html><html lang="en"> <div> <h2>AIC</h2> ...</div> <div> <p>AIC is a type of cancer that starts in the lining of the colon or rectum. ...</p> <p>AIC is a type of cancer that starts in the lining of the colon or rectum. ...</p>"	"D:\Bigdata\Medline Search Engine_Scraping_InvertedIndex\MedlineData\A_0008849.htm"

## Medline Data Snapshot in MySQL

- **Description:** This screenshot shows a preview of the data stored in the medlinedata table in MySQL. It displays multiple records, including fields such as ID, TITLE, LINK\_TO\_MEDLINE\_ARTICLE, FILEPATH, and ARTICLE\_DATA, providing an overview of the structured data available for querying.

The screenshot shows the MySQL Workbench interface with the following details:

- Navigator:** Shows the database structure with "MANAGEMENT", "INSTANCE", "PERFORMANCE", and "Administration" sections.
- Query Editor:** Displays the SQL query: `USE MedlineDB;  
select * from medlinedata;`
- Result Grid:** Shows the results of the query in a grid format with columns: ID, TITLE, LINK\_TO\_MEDLINE\_ARTICLE, FILEPATH, and ARTICLE\_DATA. The grid contains 15 rows of data.
- Output:** Shows the output of the query, which includes the count of rows (15) and the total number of rows (4483).

The screenshot shows the MySQL Workbench interface with the following details:

- Navigator:** Shows the database structure with "MANAGEMENT", "INSTANCE", "PERFORMANCE", and "Administration" sections.
- Query Editor:** Displays the SQL query: `USE MedlineDB;  
select count(*) from medlinedata;`
- Result Grid:** Shows the results of the query in a grid format with a single row labeled "count(\*)" with the value "4483".
- Output:** Shows the output of the query, which includes the count of rows (4483).

## 2. Goal of Your Intelligent Big Data Analytic Application (AI)

### Project Goal

The goal of the Medline Search Engine (MSE) project was to develop a sophisticated search platform that could provide efficient and accurate medical information to users. Specifically, the project sought to:

- **Develop a Data Collection Strategy:** Extract a large volume of medical articles from MedlinePlus.
- **Create a Data Processing Workflow:** Preprocess and clean data to make it suitable for analysis and efficient querying.
- **Build an Indexing System:** Create an inverted index to facilitate fast search queries.
- **Implementation of NLP Techniques:** Use Natural Language Processing techniques to enhance the accuracy of search results.
- **Develop a User-Friendly Search Interface:** Provide a responsive web application interface for users to enter queries and retrieve real-time search results.

### AI Components

- **Natural Language Processing (NLP):** The project leveraged various NLP techniques, such as tokenization, stemming and lemmatization to process and normalize the tokens.
- **Machine Learning Algorithms:**
  - **TF-IDF (Term Frequency-Inverse Document Frequency):** Used to measure the importance of each term in each document.
  - **Cosine Similarity:** Used to rank documents based on their relevance to user queries, providing a similarity score to determine the most relevant articles.

## 3. Platform Setting/System Configuration Procedures

### Development Environment

- **Operating System:** Windows 11
- **Programming Language:** Node.js (v20.9.0) for backend operations, JavaScript for frontend development.
- **IDE/Editor:** Visual Studio Code (VS Code) for writing and debugging code.

## System Configuration Procedures

### 1. Node.js Environment Setup:

- a. Installed Node.js and npm (Node Package Manager).
- b. Verified installations using node -v and npm -v commands.
- c. Used npm(v 10.1.0) to handle project packages.

### 2. Required Libraries Installation:

- a. Installed necessary Node.js libraries for scraping, processing, and database interactions:
  - i. **Puppeteer**: Used for web scraping (npm install puppeteer).
  - ii. **Lemmatizer, Pluralize and Stem-porter**: Libraries for NLP preprocessing (npm install lemmatizer , npm install pluralize and npm install stem-porter).
  - iii. **MongoDB and MySQL**: Drivers for database operations (npm install mongodb and npm install mysql).

### 3. MongoDB Setup:

- a. Installed MongoDB(v2.3.2) Community Edition.
- b. Created a database named MedlineDB to store raw HTML data.

### 4. MySQL Setup:

- a. Installed MySQL Server Community version (v8.0.39) for storing processed data.
- b. Created a database named MedLineDB and used MySQL Workbench for management.

### 5. Frontend Setup:

- a. Installed the Next.js framework for building a responsive web application (npx create-next-app@latest medline-search-engine).
- b. Configured necessary dependencies for creating an interactive UI.

### 6. Server Configuration:

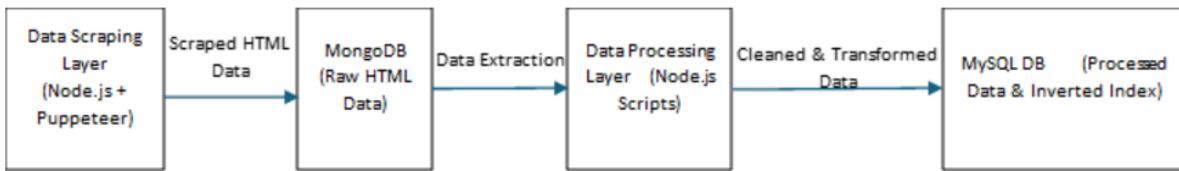
- a. Configured backend server using Node.js to handle incoming requests and process queries.Used Next.js built-in routing system.
- b. Set up local development servers for testing and debugging.

## 4. System Design (Architecture) of Your AI Application in Detail

### Overall Architecture

The system architecture of the Medline Search Engine application comprises multiple layers:

## Data Extraction and Ingestion Flow:



### 1. Data Collection Layer:

- The webScraper.js script used Puppeteer to scrape medical articles from MedlinePlus.
- Data was stored as raw HTML files and then transferred to MongoDB for backup and retrieval.

### 2. Data Processing Layer:

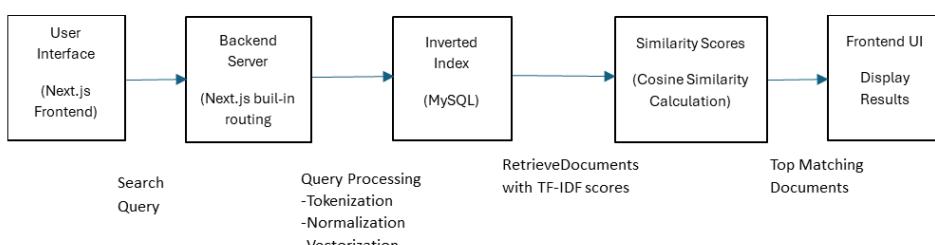
- The cleanData.js script retrieved the raw HTML content, cleaned it by removing unnecessary tags and content, and stored the processed text in articleDetails.json.
- The buildInvertIndex.js script constructed the inverted index by tokenizing articles, calculating term frequencies, and storing them in MySQL.

### 3. Data Storage Layer:

- MongoDB** was used for storing raw data scraped from the website.
- MySQL** was used for storing processed data, including document metadata and the inverted index.

### 4. Application Layer:

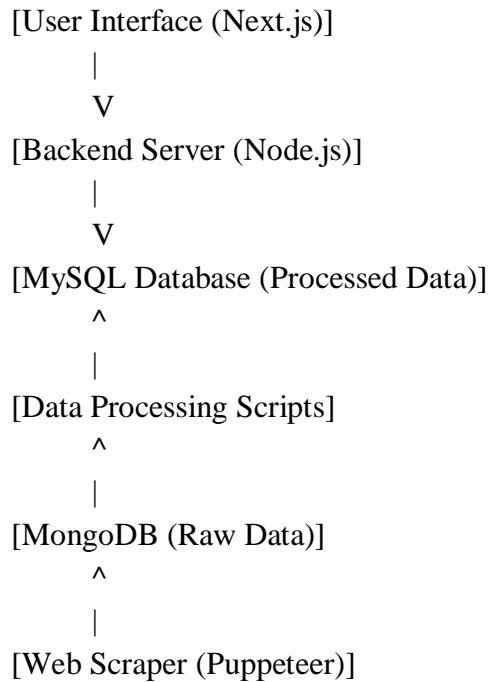
- Backend Server:** Developed with Node.js, responsible for handling requests, processing user queries, and interacting with the MySQL database.
- Frontend Interface:** Created with Next.js, providing users with a search bar to enter queries and displaying results interactively.



### 5. NLP Components:

- Implemented several NLP techniques, such as tokenization, stemming, and lemmatization.
- Used TF-IDF weighting and Cosine similarity for ranking search results and improving relevance.

## Data Flow Diagram



## 5. Raw Big Data Preprocessing Methods and Intermediate Results

### Data Cleaning (cleanData.js)

- **Extraction of Relevant Content:**
  - Identified and extracted relevant text from key HTML tags such as `<p>`, `<h1>`, `<h2>`, etc.
  - Ignored irrelevant content, including advertisements, navigation links, and redundant text.

```

import { createConnection } from "mysql2/promise";
import fs from "fs/promises";

const main = async () => {
  const conn = await createConnection({ host: "localhost", user: "root", password: "password123" });
  await createDatabase(conn, "MedLineDB");
  await createTable(conn, "MedLineData");
  // Data cleaning logic here
  conn.close();
}

```

} ;

```
await main();
```

- **Text Normalization:**

- Converted text to lowercase to ensure uniformity.
  - Removed special characters, punctuation, and unwanted symbols.
  - Addressed encoding issues and ensured the text was properly formatted for further processing.

- **Intermediate Output:**

- Cleaned articles were stored in a JSON file, articleDetails.json, containing fields like title, href, fileContent (cleaned text), and filePath.
  - This screenshot shows the articleDetails.json file containing structured data after the data cleaning process. Each record includes the article's title, href, fileContent and filePath, ready for further processing and indexing.

- Data in this format was well-structured and ready for feature extraction and indexing.

## Sample Cleaned Data Snapshot

- **Description:** This screenshot shows a sample of the cleaned content for an article. It highlights how the original HTML has been processed to extract only the meaningful text data, ensuring consistency and readability.

```
If you have cancer, a clinical trial may
If you have cancer, a clinical trial may be an option for you. A clinical trial is a study using people who agree to participate in new tests or treatments. Clinical trials help researchers know whether a new treatment works well and is safe. Trials are available for many cancers and all stages of cancer, not just advanced cancer. If you join a trial, you may get treatment that can help you. Plus, you will help others to learn more about your cancer as well as new tests or treatments. There are many things to consider before joining a trial. Learn about why you might want to enroll in a clinical trial and where to find one. What is a Clinical Trial for Cancer? Clinical trials for cancer look at ways to: Prevent cancer Screen or test for cancer Treat or manage cancer Reduce symptoms or side effects of cancer or cancer treatments A clinical trial will recruit many people to participate. During the study, each group of people will receive a different test or treatment. Some will get the new treatment being tested. Others will get standard treatment. The researchers will collect the results to see what works best. Current cancer medicines, tests, and treatments used by most health care providers have been tested through clinical trials. Should You Consider a Clinical Trial? The decision to join a clinical trial is a personal one. It is a decision you have to make based on your values, goals, and expectations. Plus, there are benefits and risks when you join a trial. Some of the benefits include: You may receive a new treatment that is not yet available to other people. You may receive treatment that is better than what is currently available. You will receive close attention and monitoring by your providers. You will help researchers understand your cancer and learn better ways to help other people with the same cancer. Some of the potential risks include: You may experience side effects. The new treatment may not work for you. The new treatment may not be as good as standard treatment. You may need more office visits and more tests. Your insurance may not pay for all of your costs in a clinical trial. Is It Safe? There are strict federal rules in place to protect your safety during a clinical trial. Safety guidelines (protocols) are agreed to before the study begins. These guidelines are reviewed by health experts to make sure that the study is based on good science and the risks are low. Clinical trials are also monitored during the entire study. Before you join a clinical trial, you will learn about the safety guidelines, what is expected of you, and how long the study will last. You will be asked to sign a consent form saying that you understand and agree to the way the study will be run and the potential side effects. Is There a Cost? Before you join a trial, make sure you look into which costs are covered. Routine cancer care costs are often covered by health insurance. You should review your policy and contact your health plan to make sure. Often, your health plan will cover most routine office visits and consults, as well as tests done to monitor your health. Research costs, such as the study medicine, or extra visits or tests, may need to be covered by the research sponsor. Also keep in mind that extra visits and tests may mean additional cost to you in lost work time and daycare or transportation costs. Who Can Join a Clinical Trial? Each clinical study has guidelines about who can join. These are called eligibility criteria. These guidelines are based on what questions the researchers are trying to answer. Studies often try to include people who have certain things in common. This can make it easier to understand the results. So you may be able to join only if you have cancer at a certain stage, are older or younger than a certain age, and do not have other health problems. If you are eligible, you can apply to be in the clinical trial. Once accepted, you become a volunteer. This means that you may quit at any time. But if you feel you want to quit, be sure you talk it over with your provider first. How to Find a Clinical Trial? Trials are done in many places, such as: Cancer centers Local hospitals Medical group offices Community clinics You can find clinical trials listed on the website of the National Cancer Institute (NCI) -- www.cancer.gov/about-cancer/treatment/clinical-trials. It is a part of the National Institutes of Health, the United States government research agency. Many of the clinical trials run across the country are sponsored by the NCI. www.cancer.gov/about-cancer/treatment/clinical-trials If you are interested in joining a clinical trial, talk with your provider. Ask if there is a trial in your area related to your cancer. Your provider can help you understand the type of care you will receive and how the trial will change or add to your care. You can also go over all of the risks and benefits to decide whether joining a trial is a good move for you. Alternative Names Intervention study - cancer References American Cancer Society website. Clinical trials. www.cancer.org/treatment/treatments-and-side-effects/
```

## 6. Design of Big Data Processing Pipeline, Data Transformation Methods

### Processing Pipeline Steps

#### 1. Data Ingestion:

- Loaded cleaned data from articleDetails.json using the uploadToDb.js script to import the content into the MySQL database.

```
import { createConnection } from "mysql2/promise";
import fs from "fs/promises";
```

```
const main = async () => {
  const conn = await createConnection({ host: "localhost", user: "root", password: "password123" });
  await createDatabase(conn, "MedLineDB");
  await createTable(conn, "MedLineData");
  // Load data into database logic here
  conn.close();
};
```

```
await main();
```

## 2. Tokenization:

- a. Split the cleaned text into individual tokens (words) using the tokenize() function to facilitate NLP processing.

## 3. Normalization:

- a. **Lowercasing:** Converted tokens to lowercase to ensure uniform representation.
- b. **Pluralization:** Reduced words to their singular form using the Pluralize library.
- c. **Stemming and Lemmatization:** Reduced words to their root form using the stem-porter and lemmatizer libraries.

## 4. Feature Extraction:

- a. Calculated **Term Frequency (TF)** to determine the number of times each term appeared in a document.
- b. Computed **Inverse Document Frequency (IDF)** to evaluate the importance of each term in the corpus.

## 5. Building the Inverted Index:

- a. Built an inverted index by mapping each term to the documents in which it appeared along with the TF-IDF scores.
- b. Stored the inverted index in MySQL for efficient retrieval and querying.

## Data Transformation Methods

- **NLP Techniques:**
  - Enhanced understanding of the text through tokenization, lemmatization and stemming.
- **Vectorization:**
  - Represented documents and queries as vectors using TF-IDF scores.
- **Indexing:**
  - Created an inverted index using MySQL to facilitate fast and efficient search operations.

## Inverted Index Table Snapshot

- **Description:** This screenshot shows a portion of the inverted\_index\_table in MySQL. It includes fields like DOC\_ID, TERM, and TERM\_FREQ, showcasing the tokenized words, their respective documents, and term frequencies.

This screenshot shows the MySQL Workbench interface with the following details:

- File Bar:** File, Edit, View, Query, Database, Server, Tools, Scripting, Help.
- Navigator:** MANAGEMENT (Server Status, Client Connections, Users and Privileges, Status and System Variables, Data Export, Data Import/Restore), INSTANCE (Startup / Shutdown, Server Logs, Options File), PERFORMANCE (Dashboard, Performance Reports, Performance Schema Setup).
- Query Editor:** Query 1 window containing the following SQL code:

```

1 USE MedlineDB;
2
3 select * from medlinedata;
4
5 select * from inverted_index_table;
6
7
8 select count(*) from medlinedata;
9
10 select * from inverted_index_table where term = "stimulating";
11
12 select count(*) from inverted_index_table;

```
- Result Grid:** Shows the data from the inverted\_index\_table. The columns are DOC\_ID, TERM, and TERM\_FREQ. The data includes rows like (1, '19', 3), (1, '2022', 3), (1, '8152022', 1), etc., up to (1, 'is', 33). A total count of 15 is shown at the bottom.
- Status Bar:** Read Only.

## Inverted Index Table Count Snapshot

- Description:** This screenshot displays the total count of entries in the inverted\_index\_table in MySQL, indicating a total of 1,421,453 indexed terms. This provides an overview of the volume of data indexed for efficient searching.

This screenshot shows the MySQL Workbench interface with the following details:

- File Bar:** File, Edit, View, Query, Database, Server, Tools, Scripting, Help.
- Navigator:** MANAGEMENT (Server Status, Client Connections, Users and Privileges, Status and System Variables, Data Export, Data Import/Restore), INSTANCE (Startup / Shutdown, Server Logs, Options File), PERFORMANCE (Dashboard, Performance Reports, Performance Schema Setup).
- Query Editor:** Query 1 window containing the following SQL code:

```

1 USE MedlineDB;
2
3 select * from medlinedata;
4
5 select count(*) from medlinedata;
6
7 select * from inverted_index_table;
8
9
10 select count(*) from inverted_index_table;
11
12 select * from inverted_index_table where term = "stimulating";
13

```
- Result Grid:** Shows the result of the count(\*) query. It displays a single row with 'count(\*)' and the value '1412453'.
- Status Bar:** Read Only.

## **7. Description of Your Knowledge Base Structure/Database Design for Information Retrieval Application**

### **Database Schema Design**

The Medline Search Engine application involved multiple databases to store different types of data, including scraped content, processed terms, and indexing tables. Here is the detailed structure:

#### **1. MongoDB (Raw Data):**

##### **a. Collection: RawHTML:**

###### **i. Fields:**

1. document\_id: Unique identifier for each document.
2. title: Title of the article.
3. url: URL of the Medline article.
4. html\_content: Original HTML content of the article.

#### **2. MySQL (Processed Data and Inverted Index):**

##### **a. Table: Documents:**

###### **i. Fields:**

1. ID: Unique identifier.
2. TITLE: Title of the document.
3. LINK\_TO\_MEDLINE: URL link.
4. ARTICLE\_DATA: Processed and cleaned content.
5. FILEPATH: Metadata information.

##### **b. Table: INVERTED\_INDEX\_TABLE:**

###### **i. Fields:**

1. DOC\_ID: Document in which term is present.
2. TERM: The word itself.
3. TERM\_FREQ: Frequency of the term in the particular document.

##### **c. Table: DICTIONARY\_TABLE:**

###### **i. Fields:**

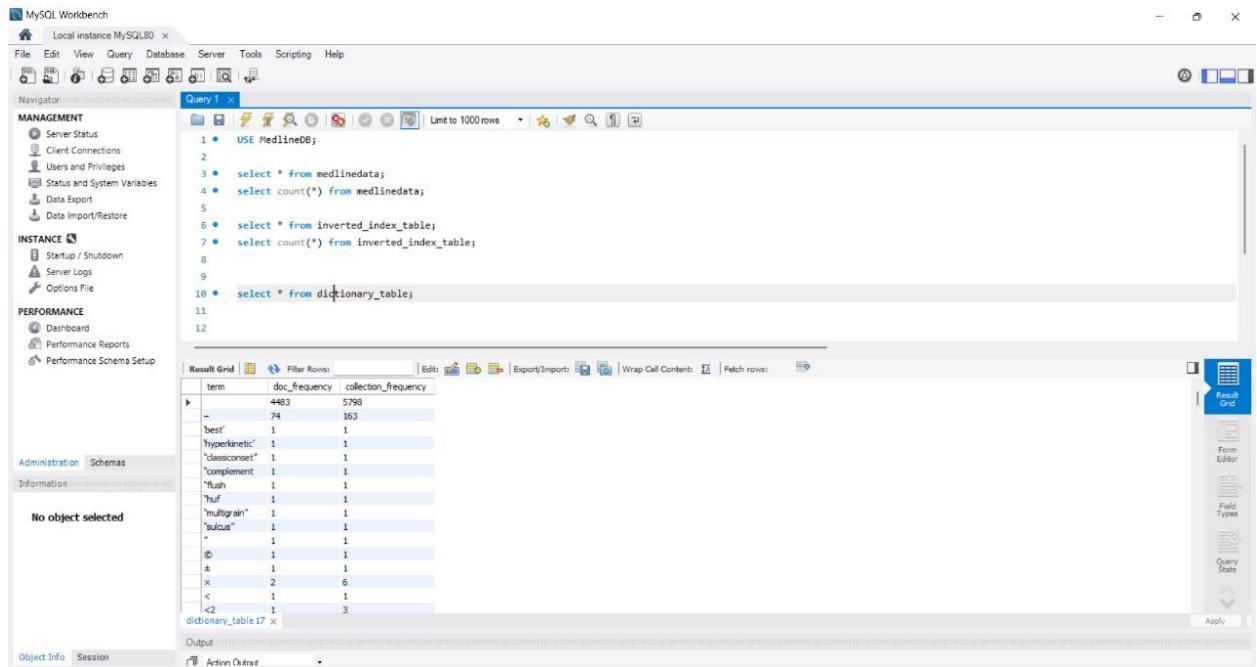
1. TERM,DOC\_FREQUENCY, TERM\_FREQUENCY: Relevant information for the mapping of terms and their frequency within documents.

## Knowledge Base Features

- **Efficient Querying:**
  - Designed with the goal of rapid information retrieval.
- **Normalization:**
  - Data normalization was done to eliminate redundancy.

## Dictionary Table Snapshot

- **Description:** Shows the DICTIONARY\_TABLE in MySQL containing key terms and their associated document frequency and collection frequency.



The screenshot shows the MySQL Workbench interface with a query editor and a results grid. The query editor contains the following SQL code:

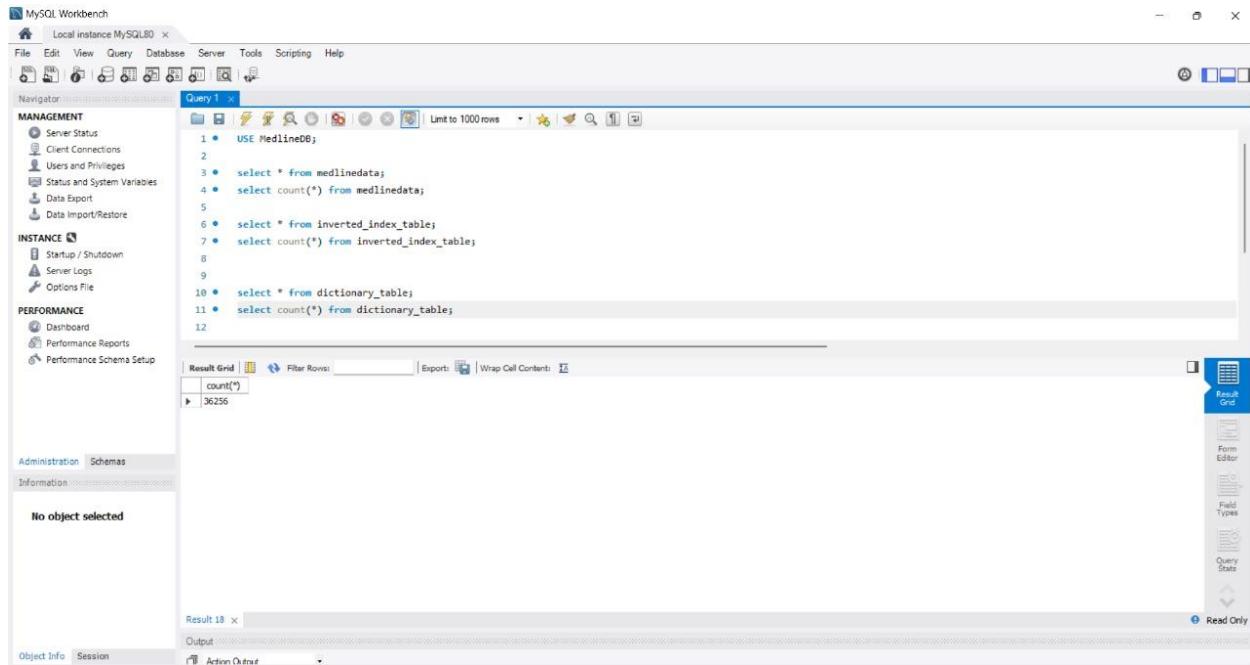
```
1 USE MedlineDB;
2
3 select * from medlinedata;
4 select count(*) from medlinedata;
5
6 select * from inverted_index_table;
7 select count(*) from inverted_index_table;
8
9
10 select * from dictionary_table;
```

The results grid displays the data from the dictionary\_table:

term	doc_frequency	collection_frequency
"	4483	5798
-	74	163
'best'	1	1
'hyperkinetic'	1	1
'classconcert'	1	1
'complement'	1	1
'flush'	1	1
'Huf'	1	1
'multigrain'	1	1
'sucus'	1	1
'"	1	1
'@'	1	1
'±'	1	1
'×'	2	6
'-	1	1
'<2'	1	3

## Dictionary Table Count Snapshot

- **Description:** Displays the total count of terms available in the dictionary\_table (36,256), indicating the scale of the data processed.



The screenshot shows the MySQL Workbench interface. In the top navigation bar, 'Query 1' is selected. The main area displays a SQL query and its results. The query is:

```
1 • USE MedlineDB;
2
3 • select * from medlinedata;
4 • select count(*) from medlinedata;
5
6 • select * from inverted_index_table;
7 • select count(*) from inverted_index_table;
8
9
10 • select * from dictionary_table;
11 • select count(*) from dictionary_table;
12
```

The result grid shows the output of the last query, which is a single row with one column:

count(*)
36256

## 8. Scoring/Ranking Algorithm

The following Information Retrieval algorithms were implemented in the project:

### 8.1 Term Frequency-Inverse Document Frequency (TF-IDF)

- **Purpose:** To evaluate the importance of a term in a document within a collection.
- **Components:**
  - **Term Frequency (TF):** Measures how frequently a term appears in a document.
  - **Inverse Document Frequency (IDF):** Reflects the importance of a term relative to the corpus size.

### 8.2 Cosine Similarity

- **Purpose:** Calculate similarity between the user's search query and the document vectors.
- **Implementation:** This was used to determine which documents were most relevant to user queries, based on their TF-IDF vectors.

## 9. The Problems/Errors Encountered and Your Resolutions

### Problems Faced

#### 1. XPath Issues During Web Scraping:

- **Problem:** During the web scraping process, the XPath used initially (articleDataXpath1 and articleDataXpath2) worked for certain elements but resulted in null data for others after an update on the website.
- **Resolution:** After extensive debugging, it was discovered that the data was located in additional classes. The XPaths were updated accordingly, which resolved the issue and enabled consistent data extraction.
- **Description:** The screenshot shows the XPaths (articleDataXpath1 and articleDataXpath2) used to identify the main content elements during scraping.

#### 2. Inefficient Database Query Execution:

- **Problem:** The initial system executed approximately 4,000 SQL queries for every user query, where each query corresponded to one document in the knowledge base. This process took around 60-75 minutes to retrieve results.
- **Resolution:** A more efficient query approach was devised. Instead of querying each document individually, a consolidated query was developed that retrieved all relevant data in a single pass. This significantly reduced the response time and improved system performance.

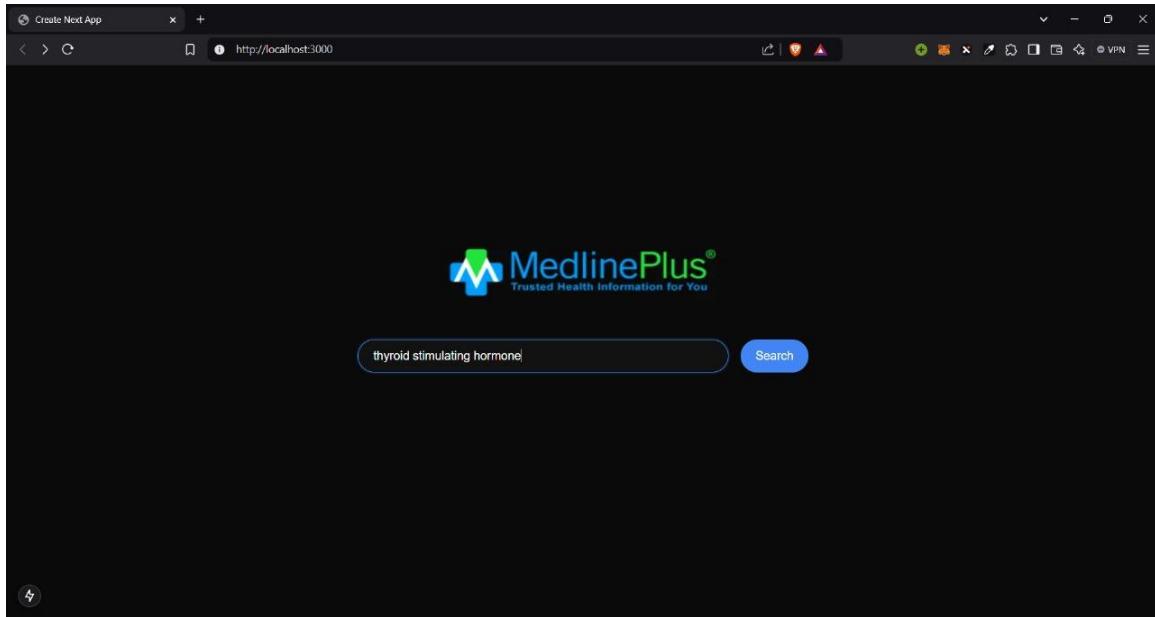
## 11. System Demo

### User Interface

- **Frontend Framework:** Built using Next.js for a responsive and interactive user interface.

### Medline UI Snapshot

- **Description:** Shows the initial interface where users can enter search queries.



## Search Demo

- **Process:**
  - Users entered a search term.
  - The backend processed the query using the inverted index and returned the top relevant results.

## Search Results Snapshot

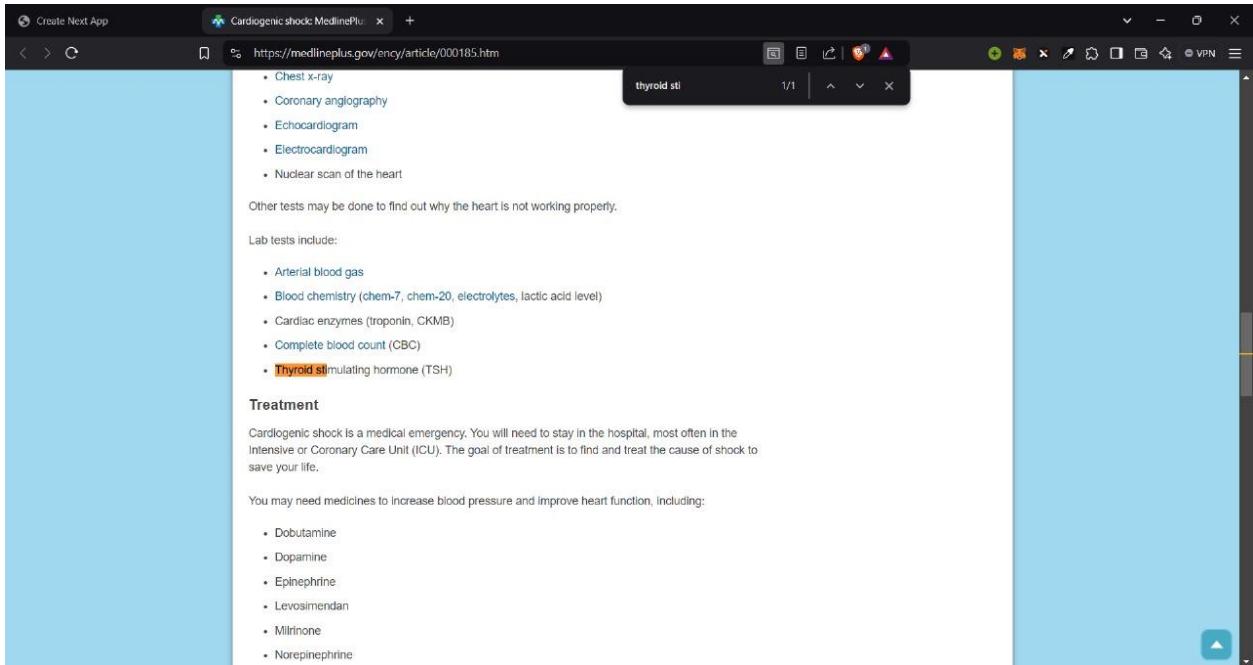
- **Description:** Demonstrates the output of a search query on the Medline UI. The results are dynamically displayed to the user with their respective titles and links.

A screenshot of a web browser window showing the search results for the query "thyroid stimulating hormone". The title "Search Results" is at the top. Below it are five search results, each in a separate box:

- Cardiogenic shock**  
<https://medlineplus.gov/ency/article/001135.htm>  
Cardiogenic shock takes place when the heart is unable to supply enough blood and oxygen to the organs of the body. Causes The most common causes of cardiogenic shock are serious heart conditions. Many of these occur during or after a heart attack (myocardial infarction). These complications include...
- Absent menstrual periods - secondary**  
<https://medlineplus.gov/ency/article/001173.htm>  
Absence of a woman's monthly menstrual period is called amenorrhea. Secondary amenorrhea is when a woman who has been having normal menstrual cycles stops getting her periods for 6 months or longer. Causes Secondary amenorrhea can occur due to natural changes in the body. For example, the most common...
- Female pattern baldness**  
<https://medlineplus.gov/ency/article/001173.htm>  
Female pattern baldness is the most common type of hair loss in women. hair loss. Causes Each strand of hair sits in a tiny hole in the skin called a follicle. In general, baldness occurs when the hair follicle shrinks over time, resulting in shorter and finer hair. Eventually, the follicle does not...
- Abnormally dark or light skin**  
<https://medlineplus.gov/ency/article/002424.htm>  
Abnormally dark or light skin is skin that has turned darker or lighter than normal. Considerations Normal skin contains cells called melanocytes. These cells produce melanin, the substance that gives skin its color. melanin. Skin with too much melanin is called hyperpigmented skin. Skin with too lit...
- Parathyroid adenoma**  
<https://medlineplus.gov/ency/article/001188.htm>  
A parathyroid adenoma is a noncancerous (benign) tumor of the parathyroid glands. The parathyroid glands are located in the neck, near or attached to the back side of the thyroid gland. benign. Causes The parathyroid glands in the neck help regulate calcium absorption, use, and removal by the body. T...

## Search Result Extended Link Snapshot

- **Description:** This shows additional information from one of the retrieved articles, providing users with detailed medical information.



## Backend Search Logic

- **Backend (route.js):**

```
import express from "express";
import mysql from "mysql2/promise";

const app = express();
const connection = await mysql.createConnection({ host: "localhost", user: "root", database: "MedLineDB" });

app.get("/search", async (req, res) => {
  const query = req.query.q;
  // Query inverted index logic here
  res.send(results);
});

app.listen(3000, () => console.log("Server running on port 3000"));
```

## Backend Code Snippet for Processing Search Queries

This code handles incoming user requests, searches the inverted index, and returns relevant documents, efficiently managing user interactions.

### Execution Steps:

#### Part 1: Backend and Data Collection Setup

1. **Unzip MedLine Search Engine Scraping InvertedIndex.zip** and navigate into it:
  - o This zip file contains all necessary backend scripts, libraries, and configurations to build the Medline Search Engine.
2. **Run npm install:**
  - o This installs all required packages and libraries (e.g., puppeteer, mysql2, mongodb).
3. **Run node webScraper.js:**
  - o This script scrapes all data from the MedlinePlus website and uploads it to MongoDB.
  - o Make sure MongoDB is installed and the MongoDB server is running on your system.
  - o **Code Overview (webScraper.js):**

```
import puppeteer from "puppeteer"
import fs from "fs"

import path from "path"
import { fileURLToPath } from "url"
import { MongoClient } from "mongodb"

const url = "https://medlineplus.gov/encyclopedia.html"
const articlesAZXPath = '//*[@id="az-section2"]//ul/li/a'
const pageArticlesXPath = '//*[@id="index"]//li/a'
const folderName = "MedlineData"
const uri = "mongodb://127.0.0.1:27017"
const dbName = "MedlineDB"
const collectionName = "MedlineData"

const writeAllArticleDetailsToFile = (articles) => {
  const filePath = "./articleHTMLFiles.json"
  const jsonData = JSON.stringify(articles, null, 4)
  fs.writeFileSync(filePath, jsonData, "utf-8")
}

const createFolder = async (folderName) => {
  // Creates a folder to store all speeches.
  try {
    fs.mkdirSync(folderName, { recursive: true }, (err) => {
      if (err) {
        console.error("Error creating folder:", err)
      } else {
        console.log(`Folder created successfully: ${folderName}`)
      }
    })
  } catch (error) {
    console.log(`Error in createFolder: ${error}`)
  }
}

const addArticleDataToFile = (anchorObj, speech) => {
  const fileName = anchorObj.title
    .replace(/\ /g, "_")
    .replace(/<>"/\/*\//g, "")
  
```

```

    .concat(".html")
  const filepath = path.join(folderName, fileName)

  fs.writeFileSync(filepath, speech)

  const fullFilePath = path.join(path.dirname(fileURLToPath(import.meta.url)), filepath)

  return fullFilePath
}

const getPageData = async () => {
  const [articalLinks, pageArticleLinks] = [[], []]
  const browser = await puppeteer.launch({
    headless: true,
  })
  const page = await browser.newPage()

  await page.goto(url, { waitUntil: "domcontentloaded" })
  const allArticles = await page.$$(`::p-xpath(${articlesAZXpath})`)

  for (let articleTag of allArticles) {
    const articleObj = await page.evaluate((el) => ({ title: el.innerHTML, href: el.href }), articleTag)
    articalLinks.push(articleObj)
  }

  for (let eachArticleLink of articalLinks) {
    await page.goto(eachArticleLink.href, { waitUntil: "domcontentloaded" })
    const allPageArticlesTags = await page.$$(`::p-xpath(${pageArticlesXpath})`)

    for (let pageArticleTag of allPageArticlesTags) {
      const pageArticleObj = await page.evaluate((el) => ({ title: el.innerHTML, href: el.href }), pageArticleTag)
      pageArticleLinks.push(pageArticleObj)
    }
  }

  for (let pageArticle of pageArticleLinks) {
    await page.goto(pageArticle.href, { waitUntil: "domcontentloaded" })
    const pageHtml = await page.content()
    pageArticle.textContent = pageHtml
    pageArticle.filePath = addArticleDatatoFile(pageArticle, pageHtml)
    console.log(`Scraped data from ${pageArticle.title}`)
  }
  writeAllArticleDetailsToFile(pageArticleLinks)
  await browser.close()
  return pageArticleLinks
}

const insertData = async (pageData) => {
  const client = new MongoClient(uri, { useNewUrlParser: true })

  try {
    await client.connect()
    console.log("Connected to MongoDB")
    const db = client.db(dbName)
    const collection = db.collection(collectionName)
    const result = await collection.insertMany(pageData)
    console.log(`${result.insertedCount} documents were inserted`)
  } catch (error) {
    console.error("Error inserting data:", error)
  } finally {
    await client.close()
  }
}

const main = async () => {
  createFolder(folderName)
  const pageData = await getPageData()
  await insertData(pageData)
}

await main()

```

- The webScraper.js uses Puppeteer to visit the MedlinePlus website, extract article titles, and save the data in MongoDB.
- writeAllArticleDetailsToFile saves the scraped data to a JSON file for backup.

#### 4. Run node cleanData.js:

- This script cleans the HTML data extracted by webScraper.js.
- It saves the cleaned data to articleDetails.json.
- **Code Overview (cleanData.js):**

```
import puppeteer from "puppeteer"

import fs from "fs/promises"

const articleDataXPath1 = '//*[@class="main"]'
const articleDataXPath2 = '//*[@class="main-single"]'

const writeAllArticleDetailsToFile = async (articles) => {
  const filePath = "./articleDetails.json"
  const jsonData = JSON.stringify(articles, null, 4)
  await fs.writeFile(filePath, jsonData, "utf-8")
}

const main = async () => {
  const browser = await puppeteer.launch()
  const page = await browser.newPage()

  const filePath = "./articleHTMLFiles.json"
  const fileContent = await fs.readFile(filePath, "utf-8")
  const pageData = JSON.parse(fileContent)
  for (let eachJsonObj of pageData) {
    const htmlString = eachJsonObj.textContent
    await page.setContent(htmlString)
    let [article] = await page.$$('p-xpath(${articleDataXPath1}))')
    if (!article) [article] = await page.$$('p-xpath(${articleDataXPath2}))')

    const articleData = await page.evaluate((element) => {
      const elements = element.querySelectorAll("p, h1, h2, h3, h4, h5, h6, li, a, b, i, u, summary")
      let.textContent = ""
      elements.forEach((el) => {
        if (el.textContent !== undefined) .textContent += el.textContent.trim() + "\n"
      })
      return.textContent
    }, article)
    eachJsonObj.textContent = articleData
  }
  writeAllArticleDetailsToFile(pageData)

  await browser.close()
}

await main()
```

- Cleans raw HTML by locating content via XPath (articleDataXPath1 and articleDataXPath2).
- Stores cleaned content into JSON for structured storage.

#### 5. Run node uploadToDb.js:

- This uploads the cleaned data to the MySQL database (MedLineDB).
- **Code Overview (uploadToDb.js):**

```
import { createConnection } from "mysql2/promise"

import fs from "fs/promises"

const host = "localhost"
const user = "root"
const password = "password123"
```

```

const dbName = "MedLineDB"
const tableName = "MedLineData"

const main = async () => {
  const conn = await createConnection({ host, user, password }) // Create a connection to MySQL server

  await createDatabase(conn, dbName)
  await createTable(conn, tableName)
  const filePath = "./articleDetails.json"
  const fileContent = await fs.readFile(filePath, "utf-8")
  const pageData = JSON.parse(fileContent)

  for (let eachJsonObj of pageData) {
    await insertIntoDB(conn, eachJsonObj)
  }

  conn.close()
}

const createDatabase = async (connection, dbName) => {
  // Creates a Database if it does not exist. dbName is already defined at top of the file
  const createDBQuery = `CREATE DATABASE IF NOT EXISTS ${dbName}`
  await connection.query(createDBQuery)
  await connection.changeUser({ database: dbName })
  console.log(`Connected to Database ${dbName}\n`)
}

const createTable = async (connection, tableName) => {
  // Creates a table if it does not exist. tableName is already defined at top of the file
  const createTableQuery = `CREATE TABLE IF NOT EXISTS ${tableName} (
    ID INTEGER PRIMARY KEY AUTO_INCREMENT,
    TITLE VARCHAR(100),
    LINK_TO_MEDLINE_ARTICLE VARCHAR(200),
    FILEPATH VARCHAR(250),
    ARTICLE_DATA MEDIUMTEXT);`
  await connection.query(createTableQuery)
  console.log(`Created Table ${tableName}\n`)
}

const insertIntoDB = async (connection, values) => {
  // Inserts the scraped data into the database
  const { title, href, filePath, fileContent } = values
  const insertDBQuery = `INSERT INTO ${tableName} (TITLE, LINK_TO_MEDLINE_ARTICLE, FILEPATH, ARTICLE_DATA) VALUES (?, ?, ?, ?)`

  await connection.execute(insertDBQuery, [title, href, filePath, fileContent])
}

await main()

```

- This script creates a MySQL table called MedLineData and inserts data into it.
- createTable ensures the table is present, while insertIntoDB adds new rows.

## 6. Run node buildInvertIndex.js:

- This script builds two tables: the dictionary and inverted index.
- **Code Overview (buildInvertIndex.js):**

```

import { createConnection } from "mysql2/promise"

import pluralize from "pluralize"
import { lemmatizer } from "lemmatizer"
import stem from "stem-porter"

export const host = "localhost"
export const user = "root"
export const password = "password123"
const createInvertedIndexTable = `CREATE TABLE IF NOT EXISTS INVERTED_INDEX_TABLE(
  DOC_ID INTEGER,
  TERM VARCHAR(250),
  TERM_FREQ INT)`

```

```

const createDictionaryTable = `CREATE TABLE DICTIONARY_TABLE (
    term VARCHAR(250) PRIMARY KEY,
    doc_frequency INT,
    collection_frequency INT
) AS
SELECT
    term,
    COUNT(DISTINCT doc_id) AS doc_frequency,
    SUM(term_freq) AS collection_frequency
FROM
    INVERTED_INDEX_TABLE
GROUP BY
    term;`

const createPostingTable = `CREATE TABLE POSTING_TABLE (
    term VARCHAR(250),
    doc_id INTEGER,
    term_freq INT
) AS
SELECT
    term,
    doc_id,
    term_freq
FROM
    INVERTED_INDEX_TABLE;`

const updateForeignKey = `ALTER TABLE POSTING_TABLE
ADD CONSTRAINT fk_term
FOREIGN KEY (term) REFERENCES DICTIONARY_TABLE(term);`


const getMedlineData = async (db) => {
    const result = await db.query(`SELECT ID, ARTICLE_DATA FROM MEDLINEDATA`)
    return result[0]
}

const createTable = async (connection, createTableQuery, tableName) => {
    // Creates a table using createTableQuery. query is defined at top of the file
    await connection.query(createTableQuery)

    console.log(`Created table ${tableName}`)
}

// Tokenizer for cleaning and processing the text

export const tokenize = (text) => {
    // Step 1: Remove punctuation
    const cleanedText = text.replace(/[\.,;—]/g, " ").replace(/\V#!$%^&&\*;{}=|+_-~()"\|\]/g, "")

    // Step 2: Convert text to lowercase
    const tokens = cleanedText.toLowerCase().split(/\s+/)

    // Step 3: Convert plurals to singular using pluralize
    const singularTokens = tokens.map((token) => pluralize.singular(token))

    // Step 4: Convert tokens to their normal form (Stemming)
    const normalizedTokens = singularTokens.map((token) => {
        return stem(token)
    })

    return normalizedTokens
}

export const getTermFrequenciesFromDoc = (tokens) => {
    const frequencies = {}

    for (const item of tokens) {
        // Increment the count if the item exists, or initialize it to 1 if it doesn't
        frequencies[item] = (frequencies[item] || 0) + 1
    }

    return frequencies
}

const insertBatchIntoDB = async (connection, rows) => {
    const insertDBQuery = "INSERT INTO INVERTED_INDEX_TABLE (TERM, DOC_ID, TERM_FREQ) VALUES ?"
    try {
        const formattedRows = rows.map(({ term, docId, frequency }) => [term, docId, frequency])
        await connection.query(insertDBQuery, [formattedRows])
    } catch (err) {
}

```

```

        console.error("Error inserting batch into DB:", err)
    }

}

const main = async () => {
    // Create a connection to MySQL server
    const medlineDB = await createConnection({ host, user, password, database: "MEDLINEDB" })

    await createTable(medlineDB, createInvertedIndexTable, "INVERTED_INDEX_TABLE")

    const allDocuments = await getMedlineData(medlineDB)

    const tokenizedDocuments = allDocuments.map((eachDocument) => {
        const id = eachDocument.ID

        return {
            id,
            tokens: tokenize(eachDocument.ARTICLE_DATA),
        }
    })

    for (const eachDocument of tokenizedDocuments) {
        const { id, title, tokens } = eachDocument
        const termFrequency = getTermFrequenciesFromDoc(tokens)

        // Prepare term frequencies for batch insertion
        const batchValues = Object.entries(termFrequency).map(([term, frequency]) => ({
            docId: id,
            term,
            frequency,
        }))
    }

    // Insert all rows for the document in a single query
    await insertBatchIntoDB(medlineDB, batchValues)

    console.log(`Inserted all terms for document ID: ${id}`)
}

await createTable(medlineDB, createDictionaryTable, "DICTIONARY_TABLE") // Subquery calculates the doc frequency and collection frequency
await createTable(medlineDB, createPostingTable, "POSTING_TABLE")
await medlineDB.query(updateForeignKey)
await medlineDB.close()
}

await main()

```

## Part 2: Frontend Setup for Search Engine UI

1. **Unzip medline-search-engine.zip** and navigate into the extracted folder:
  - o This zip file contains the frontend files needed to create the search engine web interface.
2. **Run npm install:**
  - o Installs all required packages and libraries for the Next.js application.
3. **Run npm run dev:**
  - o Starts a local development server for the web application.
4. **Navigate to http://localhost:3000:**
  - o You will see the Medline Search Engine interface where you can search for medical articles.

## TF-IDF/Cosine Similarity Logic: route.js:

```
import { createConnection } from "mysql2/promise";

import pluralize from "pluralize";
import stem from "stem-porter";
import { NextResponse } from "next/server";

const host = "localhost";
const user = "root";
const password = "password123";

// Tokenizer for cleaning and processing the text
export const tokenize = (text) => {
    // Step 1: Remove punctuation
    const cleanedText = text.replace(/\[\?!\$%\^\&\*\,\=\+\_~\)\(\]/g, "");
    // Step 2: Convert text to lowercase
    const tokens = cleanedText.toLowerCase().split(/\s+/);

    // Step 3: Convert plurals to singular using pluralize
    const singularTokens = tokens.map((token) => pluralize.singular(token));
    // Step 4: Convert tokens to their normal form (Stemming)
    const normalizedTokens = singularTokens.map((token) => {
        return stem(token);
    });
    return normalizedTokens;
};

export const getTermFrequenciesFromDoc = (tokens) => {
    const frequencies = {};

    for (const item of tokens) {
        // Increment the count if the item exists, or initialize it to 1 if it doesn't
        frequencies[item] = (frequencies[item] || 0) + 1;
    }

    return frequencies;
};

const cosineNormalize = (vector) => {
    let norm = 0;
    for (let [key, value] of Object.entries(vector)) norm += value * value;
    if (norm === 0) return vector;
    norm = Math.sqrt(norm);
    for (let eachKey of Object.keys(vector)) {
        vector[eachKey] /= norm;
    }
    return vector;
};

const cosineSimilarityScore = (queryVector, docVector) => {
    let score = 0;
    for (let key of Object.keys(queryVector)) {
        score += queryVector[key] * docVector[key];
    }

    return score;
};

const getBestMatchingDocuments = async (userQuery, medlineDB) => {
    const queryTf = getTermFrequenciesFromDoc(tokenize(userQuery));
    const uniqueTokens = Object.keys(queryTf);
    const allDocIds = (await medlineDB.query('SELECT DISTINCT(DOC_ID) FROM INVERTED_INDEX_TABLE'))[0];
    const docIds = allDocIds.map((eachDoc) => eachDoc.DOC_ID);
    const N = docIds.length;
    const queryVector = {};
    const idf = {};
    for (let token of uniqueTokens) {
        // Calculating query tf
        const frequency_raw = queryTf[token];
        const tf = 1 + Math.log10(frequency_raw);

        // Calculating idf values
        const idf_val = Math.log(N / (1 + frequency_raw));
        idf[token] = idf_val;
    }

    const scores = docIds.map((docId) => {
        let score = 0;
        for (let token of uniqueTokens) {
            const query_tf = queryVector[token];
            const doc_idf = idf[token];
            const doc_tf = docVector[token];
            const dot_product = query_tf * doc_tf * doc_idf;
            score += dot_product;
        }
        return score;
    });

    const sortedScores = scores.sort((a, b) => b - a);
    const topDocuments = docIds.slice(0, 10);
    return topDocuments;
};
```

```

const query = `SELECT DOC_FREQUENCY FROM DICTIONARY_TABLE WHERE TERM = '${token}'`;
const result = await medlineDB.query(query);
let docFrequency;
if (result[0].length == 0) {
  docFrequency = N;
} else {
  docFrequency = result[0][0]['DOC_FREQUENCY'];
}
idf[token] = Math.log10(N / docFrequency);
const tfidfWeighting = tf * idf[token];
queryVector[token] = tfidfWeighting;
}

const docVectors = {};

const queryResult = (
  await medlineDB.query(`SELECT DOC_ID, TERM, TERM_FREQ
    FROM INVERTED_INDEX_TABLE
    WHERE TERM in (${uniqueTokens.map((token) => `"' + token + "'`)}))`)
)[0];

for (let eachRecord of queryResult) {
  if (docVectors[eachRecord.DOC_ID] === undefined) {
    docVectors[eachRecord.DOC_ID] = {};
    for (let token of uniqueTokens) docVectors[eachRecord.DOC_ID][token] = 0;
  }
  docVectors[eachRecord.DOC_ID][eachRecord.TERM] = (1 + Math.log10(eachRecord.TERM_FREQ)) * idf[eachRecord.TERM];
}

const normalizedQueryVector = cosineNormalize(queryVector);
const scores = {};
for (let key of Object.keys(docVectors)) {
  docVectors[key] = cosineNormalize(docVectors[key]);
  scores[key] = cosineSimilarityScore(normalizedQueryVector, docVectors[key]);
}
const sortedScores = Object.entries(scores).sort(([valueA], [valueB]) => valueB - valueA);
// const startIndex = p * pagesPerDoc;
// const endIndex = p * pagesPerDoc + pagesPerDoc;
const results = sortedScores.map((score) => score[0]);
return results;
};

export const GET = async (params) => {
  const { url } = params;
  const searchParams = new URL(url).searchParams;
  const queryParams = {};
  for (const [key, value] of searchParams.entries()) {
    queryParams[key] = value;
  }

  const { q } = queryParams;

  try {
    const medlineDB = await createConnection({ host, user, password, database: "MEDLINEDB" });
    const results = await getBestMatchingDocuments(q, medlineDB);

    const links = (await medlineDB.query(`SELECT ID, TITLE, LINK_TO_MEDLINE_ARTICLE, ARTICLE_DATA FROM MEDLINEDATA;`))[0];

    const Objs = links.map((link) => {
      const obj = { ...link };
      obj.ARTICLE_DATA = obj.ARTICLE_DATA.slice(0, 300) + "...";
      return obj;
    });

    const sortedObjects = results.map((pId) => {
      return Objjs.find((obj) => obj.ID === pId);
    });

    await medlineDB.close();
    return NextResponse.json(sortedObjects);
  } catch (err) {
    console.log(err.message);
    return NextResponse.json({ error: err.message });
  }
}

```

};

By following these execution steps, the Medline Search Engine can be fully set up for data collection, processing, and query interfacing through the web UI. Each step has a corresponding script that has been carefully documented.

## Conclusion

The Medline Search Engine project successfully combined various big data, AI, and NLP techniques to build an efficient search platform for medical information. We utilized modern big data tools and machine learning models to build a scalable and interactive solution capable of delivering valuable medical insights to users.