# Advanced Data Structures and Algorithm Analysis

# Project 1: Roll Your Own Mini Search Engine



Date: 2024-09-30

2024-2025 Autumn&Winter Semester

# Table of Content

Chapter 1: Introduction	4
1.1 Problem Description	4
1.2 Purpose of Report	4
1.3 Backgound of Data Structures and Algorithms	4
Chapter 2: Data Structure / Algorithm Specification	5
2.1 Algorithm Architecture	5
2.2 The Main Program	6
2.3 Word Count & Stop Words	
2.4 Word Stemming	8
2.5 Inverted Index	9
2.5.1 Overall Functions	10
2.5.2 B+ Tree Operations	14
2.5.3 Hashing Operations	21
2.5.4 Other Functions	24
2.6 Query	28
2.6.1 Search Function	28
2.6.2 Search Test	30
2.6.3 Some helper functions	30
Chapter 3: Testing Results	31
3.1 Inverted Index Tests	31
3.1.1 Word Insertion Test	32
3.1.2 Single File to Multiple Files Test	38
3.1.3 Stopwords Test	41
3.1.4 Speed Test	43
3.2 Query Tests	44
3.2.1 input0	44
3.2.2 input1	45
3.2.3 input2	45
3.2.4 input3	46
3.2.5 input4	46
3.2.6 input5	47
Chapter A: Analysis and Comments	18

4.1 Space Complexity
4.2 Time Complexity
4.3 Further Improvement 51
Appendix: Source code
5.1 File Structure
5.2 getStopWord.cpp
$5.3 \text{ invIndexHeader.h} \dots 55$
5.4 invIndexFunc.cpp
5.5 invIndexTest.cpp
5.6 invIndexSearch.cpp
5.7 search_test.cpp
5.8 search_main.cpp 100
References
Declaration

# Chapter 1: Introduction

# 1.1 Problem Description

The project required us to create a **mini search engine** which can handle inquiries over "The Complete Works of William Shakespeare".

Here are some specific requirements:

- Run a word count over the Shakespeare set, extract all words from the documents by word stemming and try to identify the stop words.
- Create a customized inverted index over the Shakespeare set with word stemming. The stop words identified must not be included.
- Write a query program on top of the inverted file index, which will accept a user-specified word (or phrase) and return the IDs of the documents that contain that word.
- Run tests to show how the thresholds on query may affect the results.

# 1.2 Purpose of Report

- Show the details of the implementation of the mini search engine by show-casing essential data structures and algorithms.
- Demonstrate the correctness and efficiency of the program by analysis based on testing data and diagrams.
- Summarize the whole project, analyze the pros and cons of the mini search engine, and put forward the prospect of further improvement.

# 1.3 Backgound of Data Structures and Algorithms

- 1. **B+ Trees**: It's an improved version of search trees, widely used in the relational database and file management in operating systems. We will use this data structure to store and access to the inverted index.
- 2. **Hashing**: Hash tables have an excellent performance in searching data(only cost O(1) time), hence we take advantage of this data structure for finding stopwords when building an inverted index.

3. **Queue**: The Queue ADT is one of the most basic data structrues used in printing the B+ tree, storing the positions for terms, etc.

#### 4. **TF-IDF Algorithm**:

tf-idf (also TF\*IDF, TFIDF, TF-IDF, or Tf-idf), short for term frequency-inverse document frequency, is a measure of importance of a word to a document in a collection or corpus, adjusted for the fact that some words appear more frequently in general.

It was often used as a weighting factor in searches of information retrieval, text mining, and user modeling.

One of the simplest ranking functions is computed by summing the tf-idf for each query term; many more sophisticated ranking functions are variants of this simple model.

There are many variations of how tf-idf is calculated. In our project, we use the following formula to calculate tf, idf and tf-idf.

$$tf(word, doc) = \frac{num\ of\ word\ in\ the\ doc}{total\ word\ count\ of\ the\ doc}$$
 
$$idf(word, doc) = log\ \frac{total\ num\ of\ documents}{1 + |num\ of\ documents\ that\ include\ the\ word|}$$
 
$$ti\text{-}idf(word, doc) = tf \times idf$$

# Chapter 2: Data Structure / Algorithm Specification

# 2.1 Algorithm Architecture

The overall algorithm architecture in the program is shown below:

```
the main program

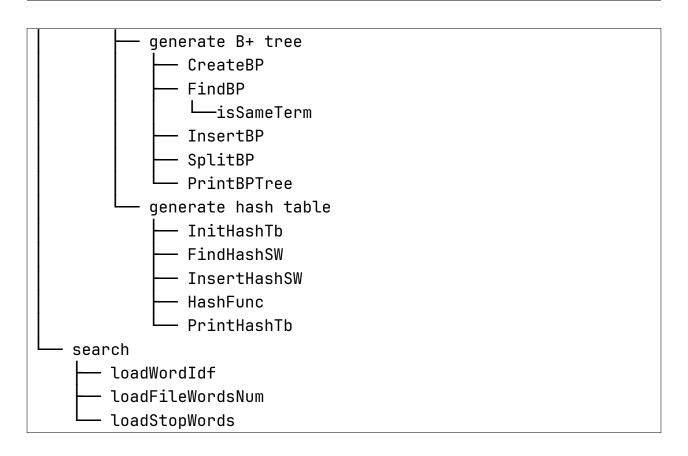
word count

stop words

inverted index

askforFilePos

fileTraversaler
```



# 2.2 The Main Program

#### Inputs: • All files of The Complete Works of Shakespeare • User's queries Outputs: • The results of queries Procedure: main() **Begin** 1 2 load necessary resources and establish inverted index 3 set values for pageSize and threshold from user input while search allow do 4 5 get query from user input 6 Calling the search function ask user if they want to continue searching or not and set 7 search\_allow accordingly end 8 9 **End**

The main idea of the code is to serve as the entry point for a search application named "ShakespeareFinder." It involves loading required resources, setting up an inverted index, and managing user queries.

# 2.3 Word Count & Stop Words

#### Note

Because the original open resources of *The Works* are written in HTML format with too many markups, we have converted them into TXT format, which is more readable and convenient for us to handle. And all titles of these articles are extracted to a TXT file at the same time.

The word count and stop words detection are relatively simple, we combined their functions into one program to cope with problems together.

**Inputs**: No obvious input parameters, but the program will read in:

- A file(code/source/txt\_title.txt) contains all titles of articles in The Works
- A directory(code/source/shakespeare\_work) includes *The Works* in TXT format

Outputs: 3 files

- code/source/stop\_words.txt: Record all selected stopwords
- code/source/word\_count.txt: Count for each word in all files
- code/source/word\_docs.txt: Count the words for each file

Procedure: getStopWord()

- 1 Begin
- 2 Read in the file txt\_title.txt as infile
- 3 Prepare the output file (outfile) named file\_word\_count.txt
- 4 while reading in the content of *infile* do
- 5 // file: one line content in infile, i.e. the title of each file
- Read in the file "shakespeare\_works/" + file + ".txt" as in
- 7 while reading each line(line) in file in do

```
8
              if find an English word (called word) then
                  Do word stemming
 9
                  wordList[word] \leftarrow wordList[word] + 1
10
                  wordNumOfDoc[file] \leftarrow wordNumOfDoc[file] + 1
11
                  wordDocs[word] \leftarrow file
12
13
              endif
14
           end
15
           Output the wordNumOfDoc to outfile
16
           closefile(in)
17
       end
       closefile(infile)
18
       Sort(wordDocs) Prepare the output file (out) named
19
       word_count.txt
20
       Prepare the output file (out2) named stop_words.txt
21
       Prepare the output file (out3) named word_docs.txt
22
       for each item(word) in wordDocs do
23
           Output the word\rightarrowcontent and word\rightarrowfrequency to out3
24
           if word \rightarrow frequency > THRESHOLD then
25
              Output the word\rightarrowcontent to out2
           endif
26
27
           Output the word \rightarrow content and wordList[word \rightarrow content] to out
28
       end
29
       closefile(out)
30
       closefile(out2)
       closefile(out3)
31
32 End
```

# 2.4 Word Stemming

We tap into the codes from a GitHub repository called "OleanderStemmingLibrary" by the author Blake-Madden. The codes are stored in the directory code/scripts/wordStem, and the link of repository is listed in the **References** section below.

#### Warning

We have to admitted that this word stemming program is kind of clumsy, especially for nouns, because the program will continue doing word stemming even though the word is in the simplest and the most common form. For example, for a simple English word "orange", it will convert it to another word "orang", which means "gorilla".

Owing to the time and capability limitation, we couldn't find a better word stemming programs in C/C++ version or convert other languages version to C/C++ version. We hope that we will use a smarter word stemming program in the foreseeable future.

#### 2.5 Inverted Index

Maybe this is the most complicated part of the whole program, because in this part we have a relatively complex algorithm architecture, and we use a couple of data structrues and algorithms, such as B+ trees, implicit queue ADT and linked list ADT. Here is the diagram of the functions used in the inverted index:

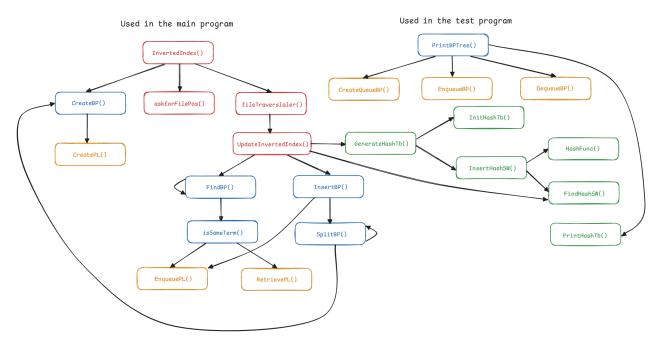


Figure 1: Relation diagram of all functions for inverted index

• Red: Overall Functions

• Blue: B+ Tree Operations

• Green: Hashing Functions

• Yellow: Other Functions

We'll introduce these functions in detail below.

#### 2.5.1 Overall Functions

#### (1) InvertedIndex

Function: The highest-level function, which users can call it directly.

```
Inputs:
• is Test: -t or --test mode, just use one particular file
• containStop Words: -s or --stopwords mode, contain stop words when
   building inverted index
Outputs:
• InvIndex: A B+ tree containing the inverted index
Procedure: InvertedIndex(is Test: bool, containStop Words: bool)
 1 Begin
 2
       InvIndex \leftarrow CreateBP()
 3
       askforFilePos(dir, fname, isTest)
 4
       // dir: directory name, fname: filename
       InvIndex \leftarrow fileTraversaler(InvIndex, dir, fname, isTest,
 5
       containStopWords)
       if InvIndex \rightarrow size > 0 then
 6
          print("Build successfully!")
 7
 8
       else
 9
          print("Fail to build an inverted index!")
```

# (2) askforFilePos

endif

return InvIndex

Function: Ask for the position of the directory or file.

#### Inputs:

10

11

12 **End** 

• *dir*: directory name

```
• fname: file name
• is Test: -t or --test mode, just use one particular file
Outputs: None, but will update either dir or fname
Procedure: askforFilePos(dir: string, fname: string, isTest: boolean)
   Begin
 1
 2
       if is Test is true then
 3
          print("Now testing the correctness of inverted Index:")
 4
          print("Please input the name of the input sample file:")
          input("Name:", fname)
 5
       else
 6
 7
           dir \leftarrow SHAKESPEAREDIR
 8
          print("Now building an inverted Index:")
          print("Do you want to search something in the default
 9
          \operatorname{path}(\{dir\})?")
          print("Please input yes or no: ")
10
11
          input(choice)
12
          switch choice
              case 'y': case 'Y':
13
14
                 break
15
              case 'n': case 'N':
16
                 print("Please input the directory of the documents:")
17
                 print("Path: ")
                 input(dir)
18
                 break
19
20
              default:
21
                 print("Choice Error!")
22
                 exit(1)
23
                 break
24
              end
          endif
25
26
       End
```

#### (3) fileTraversaler

**Function**: Make a traversal of all files(or a single file) and build the inverted index from them(or it).

```
Inputs:
• T: A B+ tree containing the inverted index
• dir: directory name
• fname: file name
• isTest: -t or --test mode, just use one particular file
• containStop Words: -s or --stopwords mode, contain stop words when
   building inverted index
Outputs: An updated B+ tree T
Procedure: fileTraversaler(T: BplusTree, dir: string, fname: string, isTest:
boolean, containStopWords: boolean)
 1 Begin
       docCnt = 0 // Count the number of documents and act as the
 2
       index of the documents at the same time
       if is Test is false then
 3
 4
          if dir exists then
              for file in the directory do
 5
 6
                 filename \leftarrow the name of file // string
                 docNames[docCnt] \leftarrow filename
 7
                 // docNames: an array containing names of
 8
                 documents(global variable)
                 wholePath \leftarrow dir + "/" + filename
 9
10
              // wholePath: the complete path of the file to be read
11
              end
12
              closefile(fp)
13
          else
14
              Error("Could not open directory!")
          endif
15
16
       else
17
           docNames[docCnt] \leftarrow fname
          dir \leftarrow DEFAULTFILEPOS // Constant: "tests" (string)
18
          wholePath \leftarrow dir + "/" + fname
19
```

```
20 endif
21 fp \leftarrow \text{openfile}(wholePath, "r") // \text{ read mode}
22 // fp: the pointer to the file
23 T \leftarrow \text{UpdateInvertedIndex}(T, docCnt, fp, containStopWords)
24 return T
25 End
```

#### (4) UpdateInvertedIndex

Function: Update the Inverted Index while reading a new document.

```
Inputs:
• T: A B+ tree containing the inverted index
• docCnt: the index of the document
• fp: pointer to the file
• containStop Words: -s or --stopwords mode, contain stop words when
   building inverted index
Outputs: An updated B+ tree T
Procedure: UpdateInvertedIndex(T: BplusTree, docCnt: integer, fp:
filePointer, containStopWords: boolean)
   Begin
 1
 2
       H \leftarrow \text{GenerateHashTb}()
 3
       while reading texts in the file pointed by fp do
 4
           if find an English word then do
 5
              term \leftarrow the English word
              if containStopWords is false and FindHashSW(term, H,
 6
              true >= 0 then
                 continue
 7
 8
              endif
 9
              term \leftarrow WordStemming(term)
10
              isDuplicated \leftarrow false
              nodebp \leftarrow FindBP(term, docCnt, T, isDuplicated)
11
12
              if isDuplicated is false then
13
                  T = \text{InsertBP}(term, docCnt, nodebp, T)
14
              endif
```

#### 2.5.2 B+ Tree Operations

#### Note

• The order of our B+ tree is 4.

#### (1) CreateBP

Function: Create a B+ tree.

```
Inputs: None
Outputs: A new and initialized B+ Tree
Procedure: CreateBP()
   Begin
 1
 2
        Allocate a memory block for new B+ tree T
 3
        for all data and children in T do
           Allocate memory blocks for term and poslist of the data, and
 4
           children
           // Use CreatePL() to intialize the poslist
 6
        end
        T \rightarrow size \leftarrow 0
        T \rightarrow childrenSize \leftarrow 0
 8
        T \rightarrow parent \leftarrow NULL
 9
10
       return T
   End
11
```

# (2) FindBP

**Function**: Find a term in B+ tree.

```
Inputs:
• term: term
   docCnt: the index of the document
  T: inverted index
• flag: true if the term is found, false otherwise
• isSearch: mark the find mode(-f or -find)
Outputs: the (possibly updated) B+ tree T or recursively call itself again
Procedure: FindBP(term: string, docCnt: integer, T: BplusTree, flag:
booleanPointer, isSearch: boolean)
 1 Begin
 2
        if T \rightarrow \text{childrenSize} = 0 then
 3
           isSameTerm(term, docCnt, T, flag, isSearch)
 4
           return T
        endif
 5
 6
        pos \leftarrow -1
        for i in range(0, T \rightarrow size) do // not contains T \rightarrow size
 7
           if term has less lexicographical order than T \rightarrow data[i] \rightarrow term
 8
           then
 9
               pos \leftarrow i
               break
10
           endif
11
12
        end
13
        if pos = -1 then
14
           pos \leftarrow i
15
        endif return FindBP(term, docCnt, T \rightarrow children[pos], isSearch)
16 End
```

# (3) isSameTerm

Function: Check if the term exists in the B+ tree.

#### Inputs:

- term: term
- docCnt: the index of the document

- nodebp: the appropriate node where the term may exists or will exists after insertion
- flag: true if the term is found, false otherwise
- *isSearch*: mark the find mode(-f or -find)

**Outputs**: None, but may update the flag and print some information regarding term

Procedure: isSameTerm(term: string, docCnt: integer, nodebp: NodeBP, flag: booleanPointer, isSearch: boolean)

```
1 Begin
 2
         if nodebp \rightarrow size > 0 then
             for i in range(0, T \rightarrow size) do // not contains T \rightarrow size
 3
                 if term = nodebp \rightarrow data[i] \rightarrow term then
 4
                     if isSearch is false then
 5
                         EnqueuePL(docCnt, nodebp \rightarrow data[i] \rightarrow poslist)
 6
 7
                     else
                         poslist \leftarrow nodebp \rightarrow data[i] \rightarrow poslist
 8
                          size \leftarrow poslist \rightarrow size
 9
                          cnt \leftarrow 0
10
11
                         print("Successfully find the word!")
12
                         print("The word was found in files below:")
13
                          posArr \leftarrow RetrievePL(poslist)
14
                         for j in range(0, size) do // not contains size
                              if posArr[j][1] <= 1 then
15
                                  print("{docNames[posArr[j][0]]}: {posArr[j][1]}
16
                                  time")
                              else
17
                                  print("{docNames[posArr[j][0]]}: {posArr[j][1]}
18
                                  times")
19
                              endif
20
                              \operatorname{cnt} \leftarrow \operatorname{cnt} + posArr|j||1|
21
                          end
22
                         print("Frequency: {cnt}")
23
                         print("----
24
                     endif
```

#### (4) InsertBP

**Function**: Insert a term into the B+ tree.

```
Inputs:
• term: term
• docCnt: the index of the document
• nodebp: the appropriate node where the term will be inserted
• Tree: B+ tree containing the inverted index
Outputs: the updated B+ tree Tree
Procedure: InsertBP(term: string, docCnt: integer, nodebp: NodeBP, Tree:
BplusTree)
1 Begin
2
       nodebp \rightarrow data[nodebp \rightarrow size] \rightarrow term \leftarrow term
       EnqueuePL(docCnt, nodebp \rightarrow data[nodebp \rightarrow size] \rightarrow poslist)
3
4
       nodebp \rightarrow size \leftarrow nodebp \rightarrow size + 1
5
       Sort(nodebp \rightarrow data)
       Tree \leftarrow SplitBP(nodebp, Tree)
6
       return Tree
   End
```

# (5) SplitBP

Function: Split the node when the node is full.

#### Inputs:

- nodebp: the appropriate node where the term will be inserted
- Tree: B+ tree containing the inverted index

```
Outputs: The updated B+ tree Tree, or recursively call itself to split
nodebp's parent node
Procedure: SplitBP(nodebp: NodeBP, Tree: BplusTree)
 1 Begin
        // ORDER: (constant)the order of B+ trees if (nodebp \rightarrow
 2
        childrenSize = 0 and nodebp \rightarrow size <= ORDER) or (nodebp \rightarrow
        childrenSize > 0 and nodebp \rightarrow size < ORDER) then
            return Tree
 3
        endif
 4
 5
 6
        // lnodebp, rnodebp: the left and right part of the split node
 7
        // tmpNodebp: store the node temporarily
 8
        // parent: the parent node of nodebp
        // cut: the position of the middle data
 9
10
11
        parent \leftarrow nodebp \rightarrow parent
12
        if parent = NULL then
13
            tmpNodebp \leftarrow CreateBP()
14
            Allocate memory for parent
15
             Tree \leftarrow parent \leftarrow tmpNodebp
16
        endif
        lnodebp \leftarrow \text{CreateBP}()
17
18
        rnodebp \leftarrow CreateBP()
19
        lnodebp \rightarrow parent \leftarrow rnodebp \rightarrow parent \leftarrow parent
20
        if nodebp \rightarrow children Size = 0 then
21
             cut \leftarrow LEAFCUT // constant: (ORDER / 2 + 1)
22
            for i in range(0, cut) do // not contains cut
23
                lnodebp \rightarrow data[i] \leftarrow nodebp \rightarrow data[i]
24
            end
25
            lnodebp \rightarrow size \leftarrow cut
            for j in range(cut, nodebp\rightarrowsize) do // not contains nodebp\rightarrow
26
            size
27
                rnodebp \rightarrow data[j - cut] \leftarrow nodebp \rightarrow data[j]
28
            end
```

```
29
               rnodebp \rightarrow size \leftarrow nodebp \rightarrow size - cut
30
          else
31
               cut \leftarrow NONLEAFCUT // constant: (ORDER / 2)
32
               for i in range(0, cut + 1) do // not contains cut + 1
33
                    if i \neq cut then
                         lnodebp \rightarrow data[i] \leftarrow nodebp \rightarrow data[i]
34
35
                    endif
                    lnodebp {\rightarrow} children[i] \leftarrow nodebp {\rightarrow} children[i]
36
37
                    lnodebp \rightarrow children \rightarrow parent \leftarrow lnodebp
38
               end
39
               lnodebp \rightarrow size \leftarrow cut
40
               lnodebp \rightarrow childrenSize \leftarrow cut + 1
               for j in range(cut + 1, nodebp \rightarrow size) do // not contains
41
               nodebp \rightarrow size
                    rnodebp \rightarrow data[j - cut - 1] \leftarrow nodebp \rightarrow data[j]
42
43
               end
               for j in range(cut + 1, nodebp \rightarrow childrenSize) do // not
44
               contains nodebp \rightarrow childrenSize
45
                    rnodebp \rightarrow children[j - cut - 1] \leftarrow nodebp \rightarrow children[j]
46
                    rnodebp \rightarrow children[j - cut - 1] \rightarrow parent \leftarrow rnodebp
47
               end
               rnodebp \rightarrow size \leftarrow nodebp \rightarrow size - cut - 1 \ rnodebp \rightarrow childrenSize
48
               \leftarrow nodebp \rightarrow childrenSize - cut - 1
49
          end
          parent \rightarrow data[parent \rightarrow size] \leftarrow nodebp \rightarrow data[cut]
50
          parent \rightarrow size \leftarrow parent \rightarrow size + 1
51
52
          if parent \rightarrow children Size > 0 then
               for i in range(0, parent \rightarrow children Size) do // not contains
53
               parent \rightarrow children Size
54
                    if parent \rightarrow children[i] = nodebp then
55
                        parent \rightarrow children[i] \leftarrow lnodebp
56
                        break
                    endif
57
               end
58
59
          else
```

```
60
               parent \rightarrow children[parent \rightarrow childrenSize] \leftarrow lnodebp
61
               parent \rightarrow childrenSize \leftarrow parent \rightarrow childrenSize + 1
62
          endif
63
          parent \rightarrow children[parent \rightarrow childrenSize] \leftarrow rnodebp
64
          parent \rightarrow childrenSize \leftarrow parent \rightarrow childrenSize + 1
65
          Sort(parent \rightarrow data)
66
          Sort(parent \rightarrow children)
67
          Tree \leftarrow SplitBP(parent, Tree)
68
          return Tree
69 End
```

#### (6) PrintBPTree

Function: Print the B+ tree(level-order traversal).

```
Inputs:
• T: B+ tree containing the inverted index
Outputs: None, but will print the whole B+ tree
Procedure: PrintBPTree(T: BplusTree)
    Begin
 1
 2
       print("B+ Tree of Inverted Index:")
 3
       q \leftarrow \text{CreateQueueBP}()
 4
       EnqueueBP(T, q)
       EnqueueBP(NULL, q)
 5
       while q \rightarrow size > 0 do
 6
 7
           nodebp \leftarrow DequeueBP(q)
 8
           if nodebp is NULL then
 9
               change to a newline
10
               if q \rightarrow size > 0 then
11
                  EnqueueBP(NULL, q)
12
               endif
13
           else
               print("[")
14
               for i in range(0, nodebp \rightarrow size) do // not contains nodebp \rightarrow
15
               size
```

```
16
                    if i = 0 then
                         print(nodebp \rightarrow data[i] \rightarrow term)
17
18
                     else
                        print(", \{nodebp \rightarrow data[i] \rightarrow term\}")
19
20
                    endif
21
                end
                print("]")
22
23
            endif
24
            if nodebp is not NULL then
                for i in range (0, nodebp \rightarrow childrenSize) do // not contains
25
                 nodebp \rightarrow children Size
                    EnqueueBP(nodebp \rightarrow children[i], q)
26
27
                end
            endif
28
29
        end
30 End
```

#### 2.5.3 Hashing Operations

# (1) GenerateHashTb

Function: Build a hash table.

```
Inputs: None
Outputs: A new hash table H, containing stopwords from the file
Procedure: GenerateHashTb()
 1
   Begin
 2
       H \leftarrow \text{InitHashTb}()
       fname \leftarrow STOPWORDPATH // constant: "stop_words.txt"
 3
       fp \leftarrow \text{openfile}(fname, "r") // \text{ read mode}
 4
       if fp is NULL then
 5
           Error("Fail to open the file of stopwords!")
 6
 7
       endif
 8
       while reading texts in the file pointed by fp do
 9
           if find an English word then do
```

```
10 \qquad term \leftarrow \text{the English word}
11 \qquad \text{InsertHashSW}(term, H)
12 \qquad \text{endif}
13 \qquad \text{end}
14 \qquad \text{closefile}(fp)
15 \qquad \text{return } H
16 \qquad \text{End}
```

#### (2) InitHashTb

Function: Initialization of the hash table.

```
Inputs: None
Outputs: A new initialized hash table
Procedure: InitHashTb()
 1
    Begin
 2
        Allocate memory block for H// HashTb
 3
        if H is NULL then
 4
            Error("Fail to create a hash table for stopwords!")
 5
        end
        H \rightarrow size \leftarrow STOPWORDSUM // maximum size
 6
 7
        for i in range(0, H \rightarrow size) do // not contains H \rightarrow size
            Allocate memory block for H \rightarrow data[i]
 8
            if H \rightarrow data[i] is NULL then
 9
10
                Error("Fail to create a hash table for stopwords!")
11
            end
12
            H \rightarrow data[i] \rightarrow stopword
            H \rightarrow data[i] \rightarrow info \leftarrow Empty // constant: 0
13
14
        end
15
        return H
16 End
```

# (3) FindHashSW

Function: Find the stopwords or other words in the hash table.

#### Inputs:

- *stopword*: stop word
- *H*: hash table containing the stop words
- justSearch: find the term without subsequent insertion

**Outputs**: A appropriate position *pos* in hash table for stopword, or just search the term in the hash table

 $\label{eq:procedure: FindHashSW (stopword: string, H: HashTb, justSearch: Procedure: FindHashSW (stopword: string, H: HashTb, FindHas$ 

```
1 Begin
```

boolean)

```
2 \qquad collisionNum \leftarrow 0
```

```
3 \quad pos \leftarrow \text{HashFunc}(stopword, H \rightarrow size)
```

if justSearch is true and  $(H \rightarrow data[pos] \rightarrow info = Empty$  or  $H \rightarrow 4$ 

```
data[pos] \rightarrow stopword = stopword) then
```

5 return -1

6 endif

while  $H \rightarrow data[pos] \rightarrow info \neq Empty$  and  $H \rightarrow data[pos] \rightarrow stopword$ 

= stopword do

 $collisionNum \leftarrow collisionNum + 1 \ pos \leftarrow pos + 2 \ *$ 

collisionNum - 1

9 if  $pos >= H \rightarrow size$  then

10  $pos \leftarrow pos - H \rightarrow size$ 

11 endif

12 **end** 

13 return pos

14 **End** 

8

# (4) InsertHashSW

Function: Insert a new stopword in hash table.

# Inputs:

- *stopword*: stop word
- $\bullet$  H: hash table containing the stop words

Outputs: None, but will update the hash table H

```
Procedure: InsertHashSW(stopword: string, H: HashTb)
   Begin
2
        pos \leftarrow \text{FindHashSW}(stopword, H, \text{false})
3
        // Legitimate: (constant) 1
        if (H \rightarrow data[pos] \rightarrow info \neq Legitimate) then
4
             H \rightarrow data[pos] \rightarrow info \leftarrow Legitimate
5
6
             H \rightarrow data[pos] \rightarrow stopword \leftarrow stopword
7
        endif
8
   \mathbf{End}
```

#### (5) HashFunc

**Function**: Hashing function.

```
Inputs:
• stopword: stop word
• size: the maximum size of the hash table
Outputs: A hash value to stopword
Procedure: HashFunc(stopword: string, size: integer)
1 Begin
2
      val \leftarrow 0
3
      for each character ch in stopword do
          val = (val << 5) + integer(ch)
4
5
      end
      return val % size
6
  \operatorname{End}
```

#### 2.5.4 Other Functions

# (1) CreateQueueBP

Function: Create the queue

```
Procedure: CreateQueueBP()

1 Begin

2 Allocate memory block for the queue Q

3 Q \rightarrow size \leftarrow 0

4 Q \rightarrow front \leftarrow Q \rightarrow rear \leftarrow 0

5 return Q

6 End
```

#### (2) EnqueueBP

**Function**: Put the node of B+ tree into the queue.

```
Inputs:
• nodebp: the newly added node
• Q: the queue
Outputs: None, but will update Q
Procedure: EnqueueBP(nodebp: NodeBP, Q: QueueBP)
1 Begin
2
       if Q \rightarrow size >= SIZE then
3
            Error("Full B+-tree-item queue!")
       endif
4
       Q \rightarrow data[Q \rightarrow rear] \leftarrow nodebp
5
       Q \!\!\to\! rear \leftarrow Q \!\!\to\! rear + 1
6
       Q \rightarrow size \leftarrow Q \rightarrow size + 1
  \operatorname{End}
```

# (3) DequeueBP

**Function**: Get the front node and delete it from the queue.

```
Inputs:
Q: the queue
Outputs: the front node returnNodeBP
Procedure: DequeueBP(Q: QueueBP)
1 Begin
```

```
2
        if Q \rightarrow size = 0 then
3
              Error("Empty B+-tree-item queue!")
4
        endif
        returnNodeBP \leftarrow Q \rightarrow data[Q \rightarrow front]
5
         Q \rightarrow front \leftarrow Q \rightarrow front + 1
6
         Q \rightarrow size \leftarrow Q \rightarrow size - 1
7
        return \ returnNodeBP
8
   End
9
```

#### (4) CreatePL

Function: Create the poslist

```
Inputs: None
Outputs: A new PosList L
Procedure: CreatePL()
1 Begin
        Allocate memory blocks for L(PosList), L \rightarrow front(PosData), L \rightarrow
2
        rear(PosData)
3
        L \rightarrow size \leftarrow 0
        L \rightarrow front \leftarrow L \rightarrow rear
4
5
        L \rightarrow rear \rightarrow pos \leftarrow -1
6
        return L
7
        End
```

#### (5) EnqueuePL

Function: Add new position.

#### Inputs:

• pos: the position

• L: the position list

Outputs: None, but will update L

Procedure: EnqueuePL(pos: integer, L: PosList)

1 Begin

```
2
           if L \rightarrow rear \rightarrow pos \neq pos then
  3
                 Allocate memory block for tmp(PosData)
  4
                 if tmp is NULL then
                      Error("Fail to create a new position data!")
  5
  6
                 endif
  7
                 tmp \rightarrow pos \leftarrow pos
 8
                 tmp \rightarrow time \leftarrow 1
                 tmp \rightarrow next \leftarrow L \rightarrow rear \rightarrow next
 9
10
                 L \rightarrow rear \rightarrow next \leftarrow tmp
11
                 L \rightarrow rear \leftarrow tmp
                 L \rightarrow size \leftarrow L \rightarrow size + 1
12
13
           else
14
                 L \rightarrow rear \rightarrow time \leftarrow L \rightarrow rear \rightarrow time + 1
15
           endif
16 End
```

#### (6) RetrievePL

Function: Retrieve all position in the list.

#### Inputs:

• L: the position list

**Outputs**: An 2D array posArr containing the all position in L, and each data contains two attributes: document index and the frequency in that document

```
Procedure: RetrievePL(L: PosList)
```

```
1
   Begin
2
        if L \rightarrow size = 0 then
3
            Error("Empty position-data queue!")
4
        endif
        Allocate memory block for posArr\ cur \leftarrow L \leftarrow front \leftarrow next
5
        i \leftarrow 0
6
7
        while cur \neq NULL do
8
            posArr[i][0] \leftarrow cur \rightarrow pos
            posArr[i][1] \leftarrow cur \rightarrow time
9
```

```
\begin{array}{lll}
10 & cur \leftarrow cur \rightarrow next \\
11 & i \leftarrow i + 1 \\
12 & end \\
13 & return posArr \\
14 & End
\end{array}
```

# 2.6 Query

#### 2.6.1 Search Function

```
Inputs:
• query: User's query in string form
 T: Inverted index
• pageSize: Number of documents to be displayed per query
• threshold: The proportion of search terms to the total number of terms
Outputs: The query result
Procedure: search(query: string, T: BplusTree, pageSize: integer,
threshold: double)
 1
   Begin
 2
       for read every character in query do
 3
          remove the Punctuation and Whitespace
 4
          if get the word then
             store the word itself and the idf of word in a vector
 5
 6
          endif
 7
       end
 8
       sort the vector in descending order
 9
       if no word then
10
          return
11
       else if one word then
12
          search the word in T
          sort the file name by tf
13
14
          print(file, tf)
15
       else
16
          according to the threshold get the word needed to searched
```

```
for every word do

search the word in T

store the result in a hash table

(the hash table store the file name and their tf-idf)

end

print(file, td-idf)

and

endif

End
```

The main idea of the function is to implement a search functionality that retrieves documents related to a given query from a B+ tree. Here is a more detailed summary of the function logic:

- 1. **Query parsing**: The code starts by loading and analyzing the query string, breaking it down into words while converting each to lowercase and stemming. It checks if each word exists in a stop words list or if it is present in the inverted index.
- 2. Valid word collection: Valid words are added to a vector, while warnings are displayed for invalid words (such as stop words or words not found in the inverted index).
- 3. **Search condition assessment**: Depending on the number of valid words, the processing is divided into two modes. If there is only one valid word, it searches directly in the inverted index and returns the matching documents with their TF values. For multiple valid words, the code calculates their TF-IDF values.
- 4. **Inverted index lookup**: For multi-word queries, the code traverses the inverted index for each valid word, using a hash table to store document IDs and their corresponding TF-IDF scores while ensuring it only retains documents that contain all the query words.
- 5. **Result sorting and output**: Finally, the results are sorted based on the TF-IDF values, and the document names along with their TF-IDF scores are printed, adhering to a specified limit on the number of results (page size).

Overall, this function provides an efficient document retrieval mechanism through valid word filtering and TF-IDF score calculation, suitable for the context of search engines or information retrieval systems.

#### 2.6.2 Search Test

```
Inputs: Files of The Works
Outputs: The results of queries
Procedure: SearchTest()
   Begin
 1
 2
       load necessary resources and establish inverted index
 3
       while test allow do
          get serial number of the test input form user
 4
 5
          get the query from the input file
 6
          get important parameters from user's input
          call the search function with the query and parameters
          ask user if they want to continue testing or not and set
 8
          test allow accordingly
 9
       end
10
  End
```

The main idea of the code is to set up a testing framework for the "Shake-speareFinder" search application. This includes loading necessary resources, allowing users to input parameters from a test file, executing search queries multiple times to evaluate performance, and managing repeated tests.

# 2.6.3 Some helper functions

# (1) Load function: loadWordIdf & loadFileWordsNum & loadStop-Words

These functions have a similar structure. The main idea is to load resources from existing files. See the pseudocode for details.

Inputs: Files of The Works
Outputs: The results of queries
Procedure: SearchTest()

- 1 Begin
- 2 load the input FILE
- 3 read resource from the file
- 4 load the resource into a hash table
- 5 End

#### (2) Search helper: FindBP2 & isSameTerm2 & RetrievePL2

These functions are very similar to their versions without the "2", but in order to better cooperate with the search function, the type of data they return is changed to vector<pair<docId, tf>>>, and the logic of getting data is changed accordingly.

# Chapter 3: Testing Results

#### 3.1 Inverted Index Tests

To verify the correctness of our inverted index, we have devised several tests from different aspects. Here is the **purpose** of each test:

- Check if every word in document(s) is inserted into the inverted index correctly.
- Build an inverted index from a single file, or a directory with a bunch of files.
- Check if the inverted index can eliminate all stopwords.

#### Warning

In the following tests, we will use the test programs to verify the correctness of our sub-programs seperately, and we won't tell you the usage of these intructions we use below. If you are curious about it, please read the README.md file in the directory code, which will guide you how to run these instructions.

#### 3.1.1 Word Insertion Test

We have two method to accomplish the first purpose: printing the whole inverted index(when the size is small), and finding the words existing in the inverted index(if words were correctly inserted).

#### (1) Printing the inverted index

```
Case 1: very simple example
$ ./invIndexTest -t -p
Now testing the correctness of inverted Index:
Please input the name of the input sample file:
Name: input1.txt
Build successfully!
B+ Tree of Inverted Index:
[beauti, ice, peach]
[appl, are, banana][beauti, cherri][ice, icecream, orang][peach,
pear, strawberri, watermelon]
input1.txt
ice
strawberry
orange
banana
peach
apple
pear
watermelon
cherry icecream you are beautiful
```

```
Case 2: simple example

$ ./invIndexTest -t -p

Now testing the correctness of inverted Index:

Please input the name of the input sample file:

Name: input2.txt

Build successfully!
```

B+ Tree of Inverted Index:

[et, lorem, nullam]

[consectetur, dolor, elit][id, ipsum][nec][pretium, sed, ut]
[adipisc, amet, at, congu][consectetur, consequat, dapibus, diam]
[dolor, e, eget][elit, erat][et, etiam, facilisi, fringilla][id,
interdum][ipsum, lacus, lectus][lorem, metus, mi][nec, nulla]
[nullam, orci, pellentesqu][pretium, purus, rhoncus][sed, sit,
sollicitudin, tincidunt][ut, vita]

input2.txt

Lorem ipsum dolor sit amet, consectetur adipiscing elit. Nullam nec erat sed nulla rhoncus dapibus et at lectus. Etiam in congue diam, ut interdum metus. Nullam pretium orci id mi pellentesque, vitae consequat lacus tincidunt. Pellentesque fringilla purus eget nulla facilisis sollicitudin.

#### (2) Finding words in the inverted index

#### Note

- This test is just used in checking the correctness of word insertion, which is similar to a simple query function, but the implementation is totally different from our formal query program, so you shouldn't mix them together.
- The texts in the following tests are too long, therefore we won't show these text in our report, but you can see them in the files positioned in the directory called code/source/tests.

#### Case 3: intermediate-level example

```
./invIndexTest -f=3
```

Now building an inverted Index:

Please input the directory of the documents:

Path: tests/input3

Build successfully!

Finding Words Mode(only supports single word finding):

Find 1: same

Successfully find the word!

The word was found in files below:

1henryiv.1.2.txt: 1 time

1henryiv.1.3.txt: 4 times

Frequency: 5

-----
Find 2: star

Successfully find the word!

The word was found in files below:

1henryiv.1.2.txt: 1 time

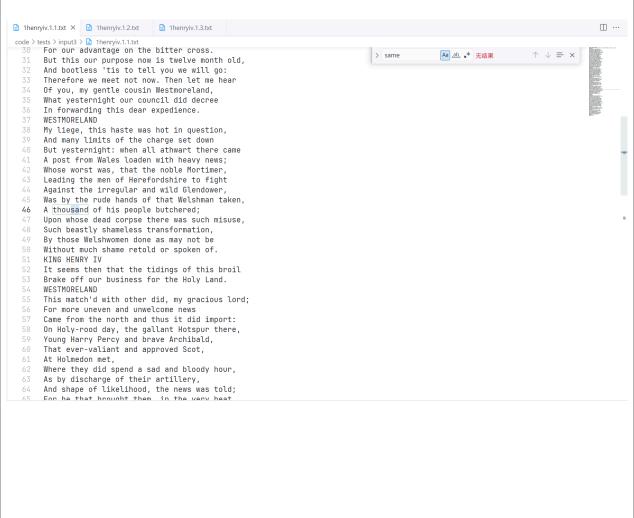
Frequency: 1

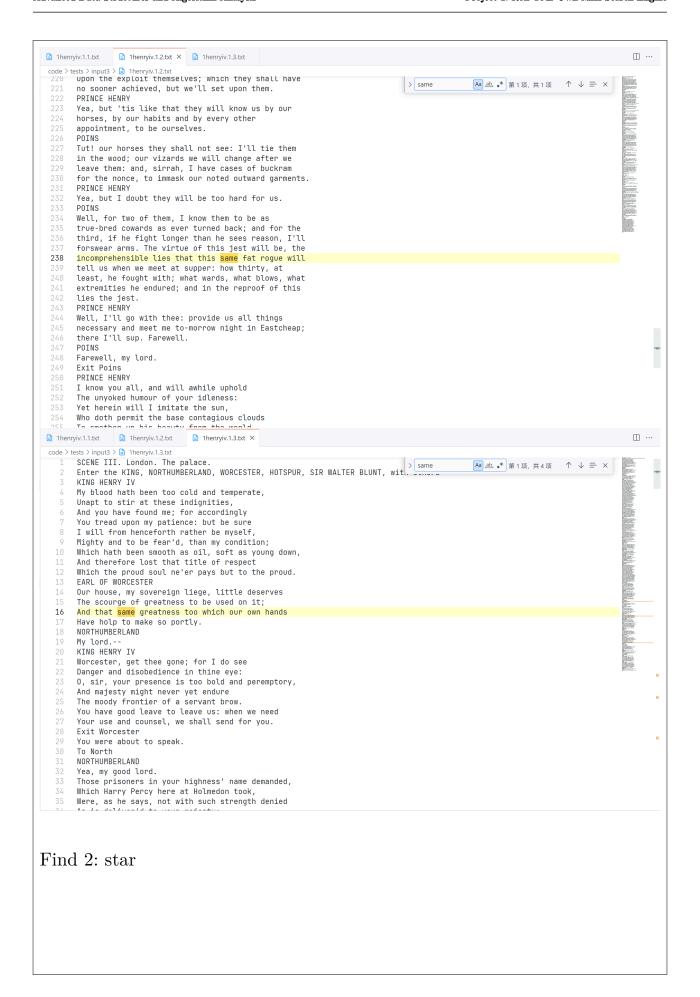
----
Find 3: fantastic

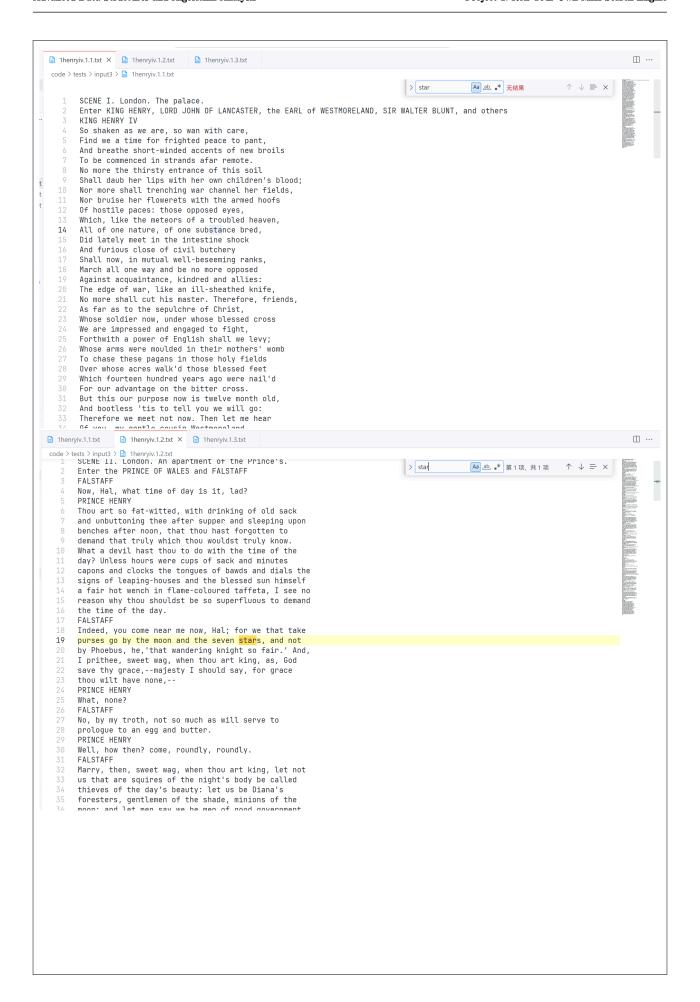
Sorry, no such word in the inverted index!

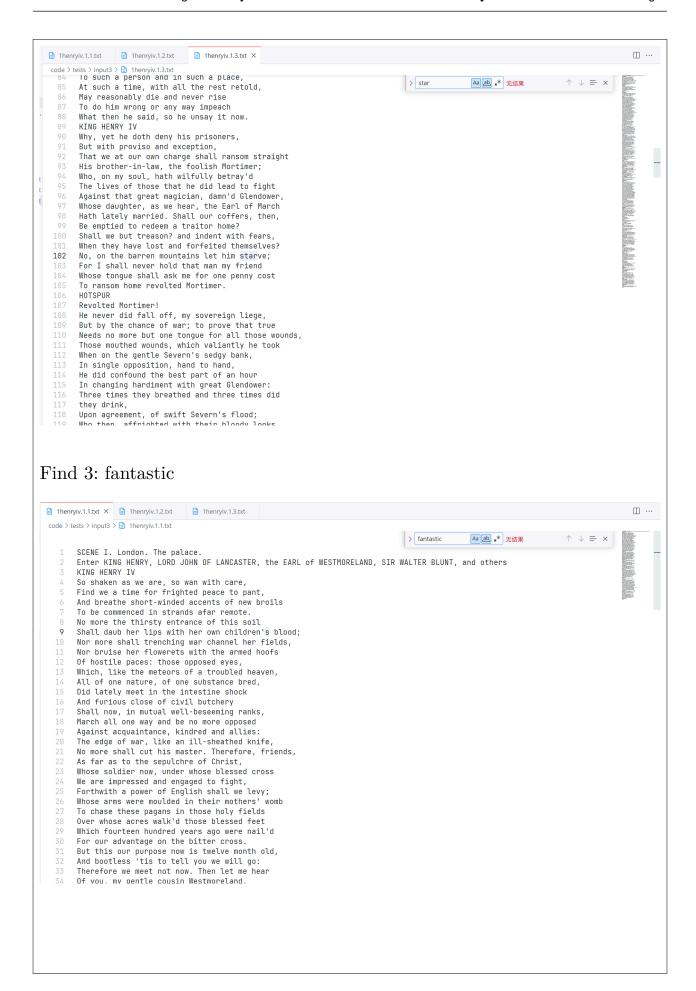
#### Verification by using finding function in Visual Studio Code

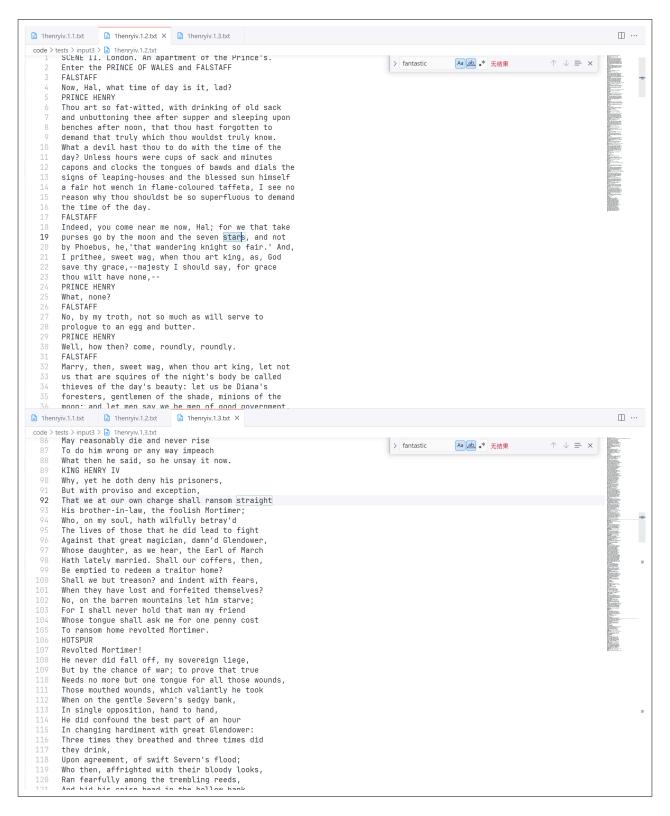
#### Find 1: same











In a nutshell, our inverted index program successfully passes the first test.

### 3.1.2 Single File to Multiple Files Test

We executed our first test based on a single file and several files, but our ultimate goal is to let our mini search engine to search something from a dozens

of files(i.e. the Complete Works of Shakespeare). So it's necessary for us to test whether the inverted index can be built from tons of files. Note that the Works is in the directory called code/source/shakespeare\_works.

Case 1: Search some dedicated words from piles of files ./invIndexTest -f=3 Now building an inverted Index: Please input the directory of the documents: Path: ../source/shakespeare\_works Build successfully! Finding Words Mode(only supports single word finding): Find 1: hamlet Successfully find the word! The word was found in files below: hamlet.1.1.txt: 3 times hamlet.1.2.txt: 42 times # Deleberate omisssion hamlet.5.1.txt: 45 times hamlet.5.2.txt: 83 times Frequency: 470 Find 2: juliet Successfully find the word! The word was found in files below: measure.1.2.txt: 3 times measure.1.4.txt: 1 time # Deleberate omisssion romeo\_juliet.5.2.txt: 1 time romeo\_juliet.5.3.txt: 19 times Frequency: 199 Find 3: macbeth Successfully find the word! The word was found in files below:

# Case 2: Search some universal words from piles of files ./invIndexTest -f=3 Now building an inverted Index: Please input the directory of the documents: Path: ../source/shakespeare\_works Build successfully! Finding Words Mode(only supports single word finding): Find 1: moon Successfully find the word! The word was found in files below: 1henryiv.1.2.txt: 5 times 1henryiv.1.3.txt: 1 time # Deleberate omisssion winters\_tale.4.3.txt: 1 time winters\_tale.4.4.txt: 1 time Frequency: 152 Find 2: happy Successfully find the word! The word was found in files below: 1henryiv.2.2.txt: 1 time 1henryiv.4.3.txt: 1 time # Deleberate omisssion winters\_tale.1.2.txt: 2 times winters\_tale.4.4.txt: 3 times

In a nutshell, our inverted index program successfully passes the second test.

# 3.1.3 Stopwords Test

Finally, we should confirm whether our program can eliminate the stopwords we have selected in advance. So we can make a comparison with two test program: one includes the stopwords, while the other doesn't include them.

```
Sorry, no such word in the inverted index!
```

```
Case 2: stopwords included
./invIndexTest -f=3 -s
Now building an inverted Index:
Please input the directory of the documents:
Path: ../source/shakespeare_works
# Delebrate omission for the display of all stopwords
Build successfully!
Finding Words Mode(only supports single word finding):
Find 1: much
# Delebrate omission for the very long position list
winters_tale.5.1.txt: 4 times
winters_tale.5.2.txt: 2 times
winters_tale.5.3.txt: 7 times
Frequency: 1070
Find 2: you
# Delebrate omission for the very long position list
winters_tale.5.1.txt: 40 times
winters_tale.5.2.txt: 19 times
winters_tale.5.3.txt: 29 times
Frequency: 14249
Find 3: great
# Delebrate omission for the very long position list
winters_tale.5.1.txt: 3 times
winters_tale.5.2.txt: 1 time
winters_tale.5.3.txt: 1 time
Frequency: 1032
```

Actually, the inverted index can eliminate all stopwords, but due to space limitation, we won't list all tests about them.

In a nutshell, our inverted index program successfully passes the third test.

Although we can't make a thorough test for the inverted index, but from the above tests, we can assure that our inverted index have no obvious error(maybe there're several small bugs existing).

### 3.1.4 Speed Test

#### Note

The specific analysis and comments about speed tests are written in **Chapter 4**.

To analyze the time complexity of the inverted index, especially the algorithms regarding the **finding** and **insertion** operations of B+ tree, we devise some timing tests for **diffrent numbers of words** in *The Works*. The results are shown below:

Number of Words(roughly)	100,000	200,000	400,000	600,000	800,000	880,000
Iterations	10	10	10	10	5	5
Ticks	1988219	3704927	7317245	11594770	7985568	8438331
Total Time(s)	1.99	3.70	7.32	11.59	7.99	8.44
Duration(s)	0.199	0.370	0.732	1.159	1.598	1.688

Table 1: Speed Tests for Inverted Index

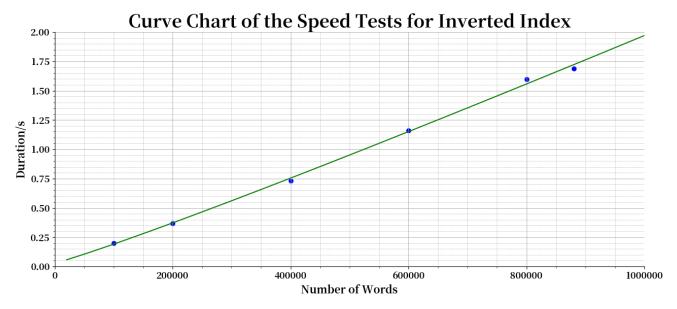


Figure 2: Curve Chart of the Speed Tests for Inverted Index

As you can see, these data points can be fitted with the product of linear function and **logarithmic function** (although the curve looks like a straight line due to the limitation of aspect ratio). So the result indicates that the time complexity of building inverted index approaches  $O(\log N)$ , and you will see the detailed explanation in Chapter 4.

# 3.2 Query Tests

Different from the oriented-procedure inverted index tests, the query tests are based on distinct and typical **input data**, which consider the performance, speed and threshold analysis simultaneously.

# 3.2.1 input0

• **content** : 1000\*'a'

• purpose: Test for illegal words of extreme length

• expected results:

Warning: "a...a" is not in the inverted index and will be ignored.

Ops, your query is in StopList or empty, so there are no documents retrieved.

• test results:

output	ticks(ticks/se	${ m earch}) \hspace{1cm} { m time}({ m s/search})$
correct	155.835	5 1.6e-4

The time is averaged from 1000 runs. Since irrelevant words will not enter the search phase, the threshold is irrelevant to this test.

# 3.2.2 input1

• **content**: peas and beans

• **purpose**: One stop word, two rare words (each appearing in only two documents), but all appearing in one document. This tests the retrieval correctness of the program and its handling of stop words.

### • expected results:

Warning: "and" is in the stop list and will be ignored.

Your query has multiple valid words, so we will search for them in the inverted index.

The words were found in files below:

File name Tf-Idf

1henryiv.2.1.txt 0.014780

#### • test result:

output	ticks(ticks/search)	${ m time}({ m s/search})$	
correct	79.439	8e-5	

The time is averaged from 1000 runs. Since the test purpose does not include the threshold, the threshold is set to 1 here.

### 3.2.3 input2

• **content** : blank input

• purpose: Testing for blank input

### expected results:

Ops, your query is in StopList or empty, so there are no documents retrieved.

#### • Test result

output	ticks(ticks/search)	$\mathrm{time}(\mathrm{s/search})$	
correct	15.246	2e-5	

The time is averaged from 1000 runs. Since irrelevant words will not enter the search phase, the threshold is irrelevant to this test.

### 3.2.4 input3

• **content** : three stop words

• **purpose**: Testing for stop words

• expected results:

Warning: "call" is in the stop list and will be ignored.

Warning: "other" is in the stop list and will be ignored.

Warning: "man" is in the stop list and will be ignored.

Ops, your query is in StopList or empty, so there are no documents

retrieved.

#### • test result:

output	${ m ticks(ticks/search)}$	$\mathrm{time}(\mathrm{s/search})$
correct	51.027	5e-5

The time is averaged from 1000 runs. Since irrelevant words will not enter the search phase, the threshold is irrelevant to this test.

# 3.2.5 input4

- **content** : Complete 1henryiv.1.2
- **purpose**: Test detection of extreme length correct text. Test the performance of searches at different thresholds
- expected results(threshold=1.0):

#### Warning:...

...(omit numorous warning about stop words)

Your query has multiple valid words, so we will search for them in the inverted index.

The words were found in files below:

File name	Tf-Idf	
1henryiv.1.2.txt	5.785513	

#### • test result:

output	ticks(ticks/search)	time(s/search)	
correct	129114.1	0.13	

The time is averaged from 10 runs, and threshold=1

${ m threshold}$	$\operatorname{rank}$	$\operatorname{tf-idf}$	$\mathbf{search} \ \mathbf{time}$
0.01	1/1	0.024940	0.02

No more thresholds were tested here because the situation was too extreme and the correct documents were already filtered out at a threshold of 0.01. Increasing the threshold will only increase the running time, so it is not tested here..

# 3.2.6 input5

- **content**: All sects, all ages smack of this vice; and he To die for't!
- **purpose**: This sentence is choose from measure.2.2. Select this to test the effect of the threshold on search results.
- expected results(threshold=1.0):

measure.2.2.txt

```
Warning: "all" is in the stop list and will be ignored.
Warning: "of" is in the stop list and will be ignored.
Warning: "of" is in the stop list and will be ignored.
Warning: "this" is in the stop list and will be ignored.
Warning: "and" is in the stop list and will be ignored.
Warning: "he" is in the stop list and will be ignored.
Warning: "to" is in the stop list and will be ignored.
Warning: "for" is in the stop list and will be ignored.
Warning: "t" is in the stop list and will be ignored.
Your query has multiple valid words, so we will search for them in the inverted index.
The words were found in files below:
File name

Tf-Idf
```

0.017464

0.015054

#### • test result:

output	${ m ticks(ticks/search)}$	time(s/search)
correct	992.903	9.9e-4

The time is averaged from 1000 runs, and threshold=1.

threshold	$\operatorname{rank}$	tf-idf	search time
0.2	3/7	0.003003	2e-4
0.4	1/1	0.005808	2e-4
0.6	1/1	0.011469	3e-4
0.8	1/1	0.012638	5e-4
1.0	1/1	0.015054	9.9e-4

From the test results, we can see that the works can be correctly screened out with a threshold of about 0.4.

I also tested some sentences with similar results, which I will not list here.

# Chapter 4: Analysis and Comments

#### Note

- We only care the memory space of major data structures
- It's undeniable that every word has different length, but we set the smallest unit to "word", not "character" for our convenience of analysis

# 4.1 Space Complexity

Conclusion: O(W + D + H + I + M + K)

- W: Word count all articles in The Works
- D: The number of documents (files, or articles in *The Works*)
- H: The size of the hash table which contains all stopwords
- I: The size of the inverted index(notice: we have removed the duplicated words, so every two nodes contain distinct words)
- M: The number of valid words
- K: The number of found documents for a single word

# Analysis:

We should analyze the space complexity step by step:

- Word count: We use C++ STL containers(pair, map and set) to store the essential information about words and files. Specifically, wordList records the word count in all files for every word; wordNumOfDoc involves the word count for every file; wordDocs contains the number of articles where words apppear for every word. Consequently, the memory space in this step depends on both word count in all files (W) and the number of files(D).
- Stop words: Actually, in the word count program, we also extracted the stop words from the variables mentioned above, then these words will be stored in a hash table for fast finding when building the inverted index. Therefore, the size of hash table(H) represents the space that stopwords use.
- Inverted index: The bulk of inverted index is stored in a B+ tree. It's universally acknowledged that the space complexity of B+ tree is O(N), when N is the number of data. However, our program has some uncertain factors, for instance, we can't control the size of the position list for each word, because the specific frequency of words are different. But what we can assure is that the total size of all position list is proportional to unduplicated word count, which is less than W. As a consequence, we only care  $I = N \times \mathsf{ORDER}$ , where  $\mathsf{ORDER}$  means the order of the B+ tree, and we allocate  $\mathsf{ORDER}$  + 1 bytes for data and children in each node.
- Query: There are four data structures used in this part: queryWord stores up to m words, yielding O(M); posVec holds document positions, with a maximum size of k, contributing O(K); freqMap may contain up to d documents, leading to O(D).; Temporary hash table currentDocIdMap also contributes O(D). Consequently, the overall space complexity is O(M + K + D).

Although our main program calls other functions which are not listed above, but their space complexity overlaps with the above operations or they are too trivial to be considered, so we don't care much about these function.

To sum up, the space complexity of our program is O(W + D + H + I + M + K).

# 4.2 Time Complexity

Conclusion:  $O(W + I \log I + n + m + m \log m + X)$ 

- W: Word count all articles in The Works
- I: The size of the inverted index
- n: The query string of length
- m: The number of valid words
- X: Vary in different situation
  - single valid word:  $k \log k$ , where k is the number of documents
  - multiple valid words:  $m \times (p \log p)$ , where p is the total document results inworst-case scenario

### **Analysis**:

- Word count and stop words: Apparently, we count and handle every word in all files to implement the functions of word count and stop words detection. Therefore, the time complexity of this part is proportional to W, which is mentioned above.
- Inverted Index: The most frequent operations we have run in the inverted index are insertion and finding, so we consider these operations mainly. It's proved that the efficiency is  $O(\log N)$  for both insertion and finding, and for every node we should execute these operations at least once. Consequently, the whole time complexity of building inverted index is  $O(I \log I)$ . Additionally, in our speed test above, we have drawn the curve chart of it, which can be fitted with  $a + b \cdot N \log N$  function, which proves the correctness of our analysis further.
- Query: We will analyze this part step by step:
  - character iteration: The function iterates through each character in the query string of length n, resulting in O(n).
  - word processing: Each word is checked against the stop words list and the inverted index. If m is the number of valid words, this step is O(m) since both checks have an average time complexity of O(1).
  - sorting: The queryWord vector containing valid words is sorted, taking  $O(m \log m)$ .
  - ocument Search: If there is one valid word, it retrieves the positions from the B+ tree (let's assume k documents) and sorts them:  $O(k \log k)$ . If there are multiple valid words, for a worst-case scenario of p total document results, it processes each word, resulting in  $O(m \times (p \log p))$ .

As a consequence:

- For a single valid word:  $O(n + m + m \log m + k \log k)$
- For multiple valid words:  $O(n + m + m \log m + m \times (p \log p))$

For the same reason above, we also ignore the time complexity of some functions which have slight impact on the whole time complexity of our main program.

In a nutshell, the time complexity of our program is  $O(W + I \log I + n + m + m \log m + X)$ .

# 4.3 Further Improvement

- 1. Algorithm refinement: So far, we have learned few of the efficient algorithms and data structures, which means that our implementation of the mini search engine might not be the best practice. However, it's possible for us to devised more ingenious and efficient procedure to cope with this problem after we systematically learned more excellent algorithms and data structures.
- 2. **Testing construction**: Although we come up with some testing cases, probably some crucial tests are still lost, and potential bugs may exists in our programs owing to our incomplete consideration. From our standpoint, it's difficult to find all typical cases for a program, but we're fully convinced that by delicate techniques and tricks for testing results, we can come up with tests as complete as possible.
- 3. Complexity analysis: As you can see, it's awkward to analyze the complexity of some programs such as the space complexity of position list in nodes of inverted index. As a consequence, our analysis on complexity isn't very accurate. We will study the systematic method of analyzing the complexity and improve the precision of our analysis in the foreseeable future.

# Appendix: Source code

# 5.1 File Structure





# 5.2 getStopWord.cpp

\* This program is used to get stop words from the Shakespeare works.\* At the same time, it also counts the number of occurrences of \* each word in each document and then gets the stop words.

\* The output files are word\_count.txt, stop\_words.txt, and word\_docs.txt.

\* All of them are stored in the code/source directory.

```
***************
#include <iostream>
#include <fstream> // for file input/output
#include <string>
#include <map>
#include <vector>
#include <set>
#include <algorithm> // for sort()
#include <cctype>
#include "wordStem/english_stem.h"
using namespace std;
typedef pair<wstring, set<string>> Pair;
map<wstring, int> wordList;
                                    // word → count(The total
number of times a word appears in the corpus)
map<string, int> wordNumOfDoc; // file → word_count(Total
word count of the file)
stemming::english_stem<> StemEnglish;
                                    // word 
ightarrow count(The
map<wstring, set<string>> wordDocs;
number of articles containing the word)
int main()
{
   string file; // file name
                     // input file
   ifstream infile;
   ofstream outfile; // output file
     infile.open("source/txt_title.txt"); // open the file
containing the file names
   outfile.open("source/file_word_count.txt",ios::out);
   while(infile >> file)
   {
       string line;
       ifstream in;
       // Read in the file and stem each word
       in.open("source/shakespeare_works/"+file+".txt", ios::in);
       while(getline(in, line))
```

```
{
            wstring word = L"";
            for(char &c : line)
            {
                // if the character is alpha or number, add it to
the word. It solve the problem of the Punctuation and Whitespace
                if(isalnum(c))
                    word += tolower(c);
                else if(word.length() > 0)
                {
                     StemEnglish(word);
                                                      //stem the
word
                    wordList[word] ++;
                    wordNumOfDoc[file] ++;
                      wordDocs[word].insert(file); //get the
number of files by the size of the file name vector
                    word = L"";
                }
            }
        outfile << file << " " << wordNumOfDoc[file] << endl;
        in.close();
    }
    infile.close();
    // Sort the words by their frequency
    // PS : Map does not have a built-in sorting algorithm, so I
need to convert the map into a vector and then use sort().
    vector<Pair> vec(wordDocs.begin(), wordDocs.end());
   sort(vec.begin(), vec.end(), [](const Pair& a, const Pair& b)
{ return a.second.size() > b.second.size(); });
    wofstream out,out2,out3; // output files
    out.open("source/word_count.txt", ios::out);
    out2.open("source/stop_words.txt", ios::out);
    out3.open("source/word_docs.txt", ios::out);
    for(Pair &word : vec)
    {
        out3 << word.first << " " << word.second.size() << endl;</pre>
```

```
// To be honest, the threshold of stop words has no
scientific basis.
        if(word.second.size() ≥ 334)
            out2 << word.first << endl; // stop words
          out << word.first << " " << wordList[word.first] <</pre>
       // word count
endl;
   out.close();
   out2.close();
   out3.close();
}
// To compile the program, run the following command in the
terminal:
// Depends on where you are, maybe you need to change the path
accordingly.
// (I assume you are in the main dictionary of the project)
// g++ -o code/scripts/getstopwords/getStopWord code/scripts/
getstopwords/getStopWord.cpp -Werror -Wall -Wextra
```

### 5.3 invIndexHeader.h

```
// Use B+ tree to store and access to the inverted index
// Declaration of properties, methods and some constants related
to B+ tree
#include <stdbool.h>
#include <stdio.h>
#include <string>
#include <time.h>
#include <vector>
#ifndef INVINDEX_H
#define INVINDEX_H // In case of re-inclusion of this header
file
#define ORDER 4
                                       // The order of B+ Tree
#define LEAFCUT (ORDER / 2 + 1)
                                        // The position of the
middle data in the leaf node of B+ Tree
#define NONLEAFCUT (ORDER / 2)
                                         // The position of the
```

```
middle data in the non-leaf node of B+ Tree
#define SIZE 1000000
                                    // The maximum size of the
queue used in printing the B+ Tree
#define MAXWORDLEN 31
                                   // The maximum length of a
single word(the longest word is about 27 or 28 in Shakespeare's
#define MAXDOCSUM 500000
                                      // The maximum number of
documents(files)
#define MAXREADSTRLEN 101
                                     // The maximum lenght of
string for one read
#define STOPWORDSUM 300
                                      // The maximum number of
stop words
of the file storing stop words
#define DEFAULTFILEPOS "source/tests"
                                             // The default
position of the file(for test mode)
#define SHAKESPEAREDIR "source/shakespeare_works" // The path
of Shakespeare's Works
                                       "source/tests/invIndex-
#define
                IISPEEDTESTDIR
speedTest/880000"
                        // Speed test files for inverted index
                    FILEWORDCOUNTPATH
#define
                                                     "source/
file_word_count.txt"
                                     // The path of word count
file
#define
                       WORDDOCSPATH
                                                     "source/
word docs.txt"
                                          // The path of word-
>doc file
#define ITERATIONS 5
                                                               //
Iteration time used in speed test for inverted index
// alias
typedef char * string;
typedef struct data * Data;
typedef struct nodebp * NodeBP;
typedef struct nodebp * BplusTree;
typedef struct poslist * Poslist;
typedef struct posdata * PosData;
typedef struct queuebp * QueueBP;
typedef struct hashtb * HashTb;
typedef struct hashsw * HashSW;
```

```
enum Kind {Legitimate, Empty}; // The state of the cells
in hash table
extern string docNames[MAXDOCSUM];  // Array containing names
of documents(global variable)
// Nodes in B+ Trees
struct nodebp {
   int size;
                                    // The size of the data in
the node
   int childrenSize;
                                    // The size of the children
nodes of the node
                                   // The data of the node
   Data data[ORDER + 1];
   NodeBP children[ORDER + 1]; // The children nodes
   NodeBP parent;
                                  // The parent node(for split
operation)
};
// Data of the node in B+ Trees
struct data {
                                   // The term
   string term;
                                    // All position where the
   PosList poslist;
term appears
};
// List of the position of terms(similar to the queue, but not
same)
struct poslist {
                                     // The size of list
   int size;
                                          // The front node of
    PosData front;
list(dummy node)
   PosData rear;
                                    // The rear node of list
};
// The specific position info
struct posdata {
   int pos;
                                    // Position, i.e. the index
of the document
```

```
int time;
                                    // the frequency in a single
document
                                     // the next pointer
   PosData next;
};
// The queue of nodes in B+ tree(array implementation)
struct queuebp {
                                      // The current size of queue
   int size;
   int front;
                                      // The index of the front
node
                                       // The index of the rear
    int rear;
node
                                     // Data
   NodeBP data[SIZE];
};
// The hash table for stop words
struct hashtb {
                                     // The maximum size of the
   int size;
hash table
   HashSW data[STOPWORDSUM];
                              // Data
};
// The cells in hash table
struct hashsw {
   string stopword;
                                     // Stop word
                                     // State. either legitimate
   enum Kind info;
or empty
};
// All methods are listed here. The explanation of parameters are
in the file "invIndexFunc.cpp"
// Methods for building inverted index
// The highest-level function, which users can call it directly
            InvertedIndex(bool
                                  isTest = false, bool
BplusTree
containStopWords = false);
// Ask for the position of the directory or file
void askforFilePos(char * dir, char * fname, bool isTest);
// Make a traversal of all files(or a single file) and build the
inverted index from them(or it)
```

```
BplusTree fileTraversaler(BplusTree T, char * dir, char * fname,
bool isTest, bool containStopWords);
// Update the Inverted Index while reading a new document
BplusTree UpdateInvertedIndex(BplusTree T, int docCnt, FILE * fp,
bool containStopWords);
// Methods about B+ tree
// Create a B+ tree
BplusTree CreateBP();
// Find a term in B+ tree
NodeBP FindBP(string term, int docCnt, BplusTree T, bool * flag,
bool isSearch = false);
// Check if the term is in the B+ tree
void isSameTerm(string term, int docCnt, NodeBP nodebp, bool *
flag, bool isSearch = false);
// Insert a term into the B+ tree
BplusTree InsertBP(string term, int docCnt, NodeBP
                                                         nodebp,
BplusTree Tree);
// Split the node when the node is full
BplusTree SplitBP(NodeBP nodebp, BplusTree Tree);
// Print the B+ tree(level-order traversal)
void PrintBPTree(BplusTree T);
// Methods about the queue
// Create the queue
QueueBP CreateQueueBP();
// Put the node of B+ tree into the queue
void EnqueueBP(NodeBP nodebp, QueueBP Q);
// Get the front node and delete it from the queue
NodeBP DequeueBP(QueueBP Q);
// Methods about poslist
// Create the poslist
PosList CreatePL();
// Add new position
void EnqueuePL(int pos, PosList L);
// Retrieve all position in the list
int ** RetrievePL(PosList L);
```

```
// Methods about hash table
// Build a hash table
HashTb GenerateHashTb();
// Initialization of the hash table
HashTb InitHashTb();
// Find the stopwords or other words in the hash table
int FindHashSW(string stopword, HashTb H, bool justSearch);
// Insert a new stopword in hash table
void InsertHashSW(string stopword, HashTb H);
// Hashing function
int HashFunc(string stopword, int size);
// Print hash table
void PrintHashTb(HashTb H);
// Comparison functions used in qsort()
int cmpData(const void * a, const void * b);  // Compare data
of the node in B+ tree
int cmpNodeBP(const void * a, const void * b); // Compare the
node by their data
// wstring \longleftrightarrow char *, for word stemming
std::wstring chararrToWstring(char * st);
char * wstringToChararr(std::wstring wst);
// Word Stmming wrapper
string WordStem(string term);
// Print the ticks and duration, for -tr or --time function
void PrintTime(clock_t start, clock_t end);
void loadWordIdf(std::string filePath);
void loadFileWordsNum(std::string filePath);
void loadStopWords(std::string filePath);
void search(std::string query, BplusTree T, int pageSize, double
threshold);
std::vector<std::pair<int,double>> FindBP2(string term,
                                                              int
docCnt, BplusTree T) ;
std::vector<std::pair<int,double>> isSameTerm2(string term, int
docCnt, NodeBP nodebp) ;
```

```
std::vector<std::pair<int,double>> RetrievePL2(PosList L) ;
#endif
```

# 5.4 invIndexFunc.cpp

```
// Implementation of methods related to B+ tree in invIndex.h
#include "invIndexHeader.h"
#include "wordStem/english_stem.h"
#include <algorithm>
#include <codecvt>
#include <filesystem>
#include <locale>
#include <string>
// To avoid unexpected import problems
extern "C" {
   #include <ctype.h>
   #include <stdbool.h>
   #include <stdio.h>
   #include <stdlib.h>
   #include <string.h>
}
// file system namespace
namespace fs = std::filesystem;
string docNames[MAXDOCSUM]; // Array containing names of
documents(global variable)
HashTb H;
                                 // Hash table storing the stop
words
// The highest-level function, which users can call it directly
// isTest: -t or --test mode, just use one particular file
// containStopWords: -s or --stopwords mode, contain stop words
when building inverted index
BplusTree InvertedIndex(bool isTest, bool containStopWords) {
                                        // Directory name
   char dir[MAXREADSTRLEN];
                                        // File name
   char fname[MAXREADSTRLEN];
   BplusTree InvIndex = CreateBP();  // Inverted index, stored
in B+ tree
```

```
// askforFilePos(dir, fname, isTest); // Ask for the position
of file or directory
    // Make a traversal in the directory(or a single file) and
build the inverted index from it
    strcpy(dir, SHAKESPEAREDIR);
   isTest = false;
     InvIndex = fileTraversaler(InvIndex, dir, fname, isTest,
containStopWords);
   if (InvIndex→size) {
                                       // If the inverted index
contains the data, it indicates the success of building
        printf("Build successfully!\n");
   } else {
                                         // Otherwise, it fails
        printf("Fail to build an inverted index!\n");
   }
                                           // Return the final
    return InvIndex;
inverted index
}
// Ask for the position of the directory or file
// dir: directory name
// fname: file name
// isTest: -t or --test mode, just use one particular file
void askforFilePos(char * dir, char * fname, bool isTest) {
    if (isTest) {     // If we choose the test mode in the test
file,
        printf("Now testing the correctness of inverted Index:
\n");
        printf("Please input the name of the input sample file:
\nName: ");
        scanf("%s", fname); // we should input the filename
    } else { // Otherwise(in the main program or other
default situations),
        char choice[MAXWORDLEN];
        strcpy(dir, SHAKESPEAREDIR); // Default path
        printf("Now building an inverted Index:\n");
        printf("Do you want to search something in the default
path(%s)?\n", dir);
```

```
printf("Please input yes or no: ");
                               // Input your choice
       scanf("%s", choice);
       switch (choice[0]) {
            case 'y': case 'Y': // Yes
               break:
           case 'n': case 'N':
                                    // No
                   printf("\nPlease input the directory of the
documents:\nPath: ");
               scanf("%s", dir); // We should input the name
of the directory
               break;
            default:
                                     // Choice error
               printf("Choice Error!\n");
               exit(1);
               break;
       }
   }
}
// Make a traversal of all files(or a single file) and build the
inverted index from them(or it)
// T: B+ tree containing the inverted index
// dir: directory name
// fname: file name
// isTest: -t or --test mode, just use one particular file
// containStopWords: -s or --stopwords mode, contain stop words
when building inverted index
BplusTree fileTraversaler(BplusTree T, char * dir, char * fname,
bool isTest, bool containStopWords) {
   int docCnt = 0:
                       // Count the number of documents and act
as the index of the documents at the same time
   char * wholePath; // The whole path name
   FILE * fp = NULL; // File pointer
   H = GenerateHashTb();
                                         // Build a hash table
for stop words
    if (containStopWords) {
                                            // If in stopwords
mode, print the hash table
       PrintHashTb(H);
```

```
}
   wholePath = new char[MAXREADSTRLEN];
    if (!isTest) {      // If we choose the test mode in the
test file,
        fs::path dirPath(dir);
          if (fs::exists(dirPath) && fs::is_directory(dirPath))
{
   // Make a traversal in the directory
         for (const auto& entry : fs::directory_iterator(dirPath))
{
               if (fs::is_regular_file(entry)) {  // entry: a
single file
                                       std::string filename =
entry.path().filename().string(); // Get the file name
                   docNames[docCnt] = new char[filename.length()
+ 1];
                 strcpy(docNames[docCnt], filename.c_str()); //
Store the filename
                    strcpy(wholePath, (dirPath.string() + "/" +
filename).c_str()); // Get the whole path name
                    // Open the file
                    fp = fopen(wholePath, "r");
                    if (!fp) { // Error handler
                        printf("Couldn't open the file!\n");
                        exit(1);
                    }
                    // Update the inverted index
                       T = UpdateInvertedIndex(T, docCnt++, fp,
containStopWords);
            if (fp) // Don't forget close the file pointer
               fclose(fp);
        } else { // Input wrong directory
           perror("Could not open directory");
        }
   } else {
       strcpy(dir, DEFAULTFILEPOS); // The file is in the
```

```
default position
        std::string sdir(dir);
        std::string sfname(fname);
       strcpy(wholePath, (sdir + "/" + sfname).c_str()); // Get
the whole path name
             docNames[docCnt] = (string)malloc(sizeof(char) *
(strlen(fname) + 1));
        strcpy(docNames[docCnt], fname);
                                                              //
Store the filename
        // Open the file
        fp = fopen(wholePath, "r");
        if (!fp) { // Error handler
            printf("Couldn't open the file!\n");
            exit(1);
        }
        // Update the inverted index
                     = UpdateInvertedIndex(T, docCnt++, fp,
containStopWords);
        fclose(fp); // Don't forget close the file pointer
    }
    return T;
}
// Update the Inverted Index while reading a new document
// T: B+ tree containing the inverted index
// docCnt: the index of the document
// fp: file pointer
// containStopWords: -s or --stopwords mode, contain stop words
when building inverted index
BplusTree UpdateInvertedIndex(BplusTree T, int docCnt, FILE * fp,
bool containStopWords) {
    int i;
   int pre, cur;
                                          // Mark the start and
the end of one word
   char tmp[MAXREADSTRLEN];
                                         // Memory space storing
the reading data temporarily
                                           // Term(or word)
    string term;
```

```
bool isDuplicated;
                                              // A flag, record
whether the term exists in the B+ tree
                                          // Node in B+ tree
   NodeBP nodebp;
     while (fgets(tmp, MAXREADSTRLEN - 1, fp) ≠ NULL) { //
Continue reading the file, until arrive at the end of file
        pre = cur = 0;
                                             // Initialization
        for (i = 0; i < strlen(tmp); i++) { // Retrieve all</pre>
characters in the tmp string
           if (!isalpha(tmp[i])) {  // Maybe it's time
to record a word
                cur = i;
                  if (cur > pre) {
                                                  // Legitimate
situation
                    term = (char *)malloc(sizeof(char) * (cur -
pre + 1));
                   strncpy(term, tmp + pre, cur - pre);
                     term[cur - pre] = '\0'; // Don't forget
this step
                   // Word stemming
                   term = WordStem(term);
                   // If we consider the stop words(default) and
assure the term is a stop word,
                   if (!containStopWords && FindHashSW(term, H,
true) ≥ 0) {
                       pre = cur + 1;
                       continue; // then we should ignore it
                   }
                   isDuplicated = false;
               nodebp = FindBP(term, docCnt, T, &isDuplicated);
Find the appropriate position for the term
                   // If isDuplicated is true, then the time of
the term will +1 in function isSameTerm()
                   if (!isDuplicated) { // If it's a new term,
insert it!
```

```
T = InsertBP(term, docCnt, nodebp, T);
                    }
                }
               pre = cur + 1;  // Move the start position for
possible new word
        }
        // Handle the last possible word in the tmp string
        if (!cur || pre > cur \&\& pre \neq i) {
            cur = i;
             term = (char *)malloc(sizeof(char) * (cur - pre +
1));
            strncpy(term, tmp + pre, cur - pre);
            term[cur - pre] = '\0';
            // Word stemming
            term = WordStem(term);
            // If we consider the stop words(default) and assure
the term is a stop word,
           if (!containStopWords && FindHashSW(term, H, true) ≥
0) {
                pre = cur + 1;
                continue; // then we should ignore it
            }
            isDuplicated = false;
            nodebp = FindBP(term, docCnt, T, &isDuplicated); //
Find the appropriate position for the term
           // If isDuplicated is true, then the time of the term
will +1 in function isSameTerm()
            if (!isDuplicated) { // If it's a new term, insert
it!
                T = InsertBP(term, docCnt, nodebp, T);
            }
        }
    }
```

```
return T;
}
// Create a B+ tree
BplusTree CreateBP() {
    BplusTree T = (BplusTree)malloc(sizeof(struct nodebp)); //
Allocate the memory space for new B+ tree
    if (T = NULL) { // Allocation failure
        printf("Failed to create a B+ Tree!\n");
        return T;
    }
    int i;
    // Memory allocation and initialization of data and children
    for (i = 0; i \leq ORDER; i \leftrightarrow) \{
        T→data[i] = (Data)malloc(sizeof(struct data));
              T→data[i]→term = (string)malloc(sizeof(char) *
MAXWORDLEN);
        T \rightarrow data[i] \rightarrow poslist = CreatePL();
        T→children[i] = (NodeBP)malloc(sizeof(struct nodebp));
    }
    // Initialization of other fields
    T \rightarrow size = 0;
    T→childrenSize = 0;
    T \rightarrow parent = NULL;
    return T;
}
// Find a term in B+ tree
// term: term
// docCnt: the index of the document
// T: inverted index
// flag: true if the term is found, false otherwise
// isSearch: mark the find mode(-f or --find)
NodeBP FindBP(string term, int docCnt, BplusTree T, bool * flag,
bool isSearch) {
```

```
int i;
   if (!T) { // If the tree is empty, return the tree(actually,
it's impossible in our program)
        return T:
    } else if (!T→childrenSize) { // If we arrive at the leaf
node, search its data
        isSameTerm(term, docCnt, T, flag, isSearch);
        return T;
    }
    int pos = -1; // The index of the appopriate non-leaf node
    for (i = 0; i < T \rightarrow size; i \leftrightarrow) {
         if (strcmp(term, T \rightarrow data[i] \rightarrow term) < 0) { // Find the}
first node which have term with higher lexicographic number
            pos = i;
            break;
        }
    }
     if (pos = -1) { // If no position found in above loop,
choose the last node
        pos = i;
    }
        return FindBP(term, docCnt, T→children[pos],
isSearch); // Continue finding in the children node
}
// Check if the term exists in the B+ tree
// term: term
// docCnt: the index of the document
// nodebp: the appropriate node where the term may exists or will
exists after insertion
// flag: true if the term is found, false otherwise
// isSearch: mark the find mode(-f or --find)
void isSameTerm(string term, int docCnt, NodeBP nodebp, bool *
flag, bool isSearch) {
    int i;
```

```
if (nodebp→size) { // If it's not an empty node, start
searching
        for (i = 0; i < nodebp \rightarrow size; i ++) {
           if (!strcmp(term, nodebp\rightarrowdata[i]\rightarrowterm)) { // If
the term exists in the inverted index
                if (!isSearch) {
                                      // If it's not in the find
mode
                  EnqueuePL(docCnt, nodebp→data[i]→poslist); //
Update the poslist of the term
                } else { // Otherwise, print all info of the
term
                 PosList poslist = nodebp→data[i]→poslist;
Position list
                  int size = poslist→size;
The number of all documents where the term appears
                   int cnt = 0;
Record the total frequency of the term
                 printf("Successfully find the word!\n");
Some banners
                     printf("The word was found in files below:
\n");
                    int j;
                    int ** posArr = (int **)malloc(sizeof(int *)
* size); // Allocation of a 2D array
                    for (j = 0; j < size; j++) {
                         posArr[i] = (int *)malloc(sizeof(int) *
2);
                    }
                  posArr = RetrievePL(poslist);
Put the poslist in a 2D array
                     for (j = 0; j < size; j++) \{ // Print the \}
name of documents and their frequency respectively
                        if (posArr[j][1] ≤ 1) // Singular
                         printf("%s: %d time\n", docNames[posArr[j]]
[0]], posArr[j][1]);
```

```
else // Plural
                       printf("%s: %d times\n", docNames[posArr[j]]
[0]], posArr[j][1]);
                       cnt += posArr[j][1];
                 printf("Frequency: %d\n", cnt);
The total frequency
                    printf("-----
\n");
                }
                *flag = true; // mark the flag, indicating
we find the term
                break;
            }
        }
   }
}
// Insert a term into the B+ tree
// term: term
// docCnt: the index of the document
// nodebp: the appropriate node where the term will be inserted
// Tree: B+ tree containing the inverted index
BplusTree InsertBP(string term, int docCnt, NodeBP nodebp,
BplusTree Tree) {
   int i;
    strcpy(nodebp→data[nodebp→size]→term, term);
   EnqueuePL(docCnt, nodebp\rightarrowdata[nodebp\rightarrowsize++]\rightarrowposlist); //
Add the data info
     qsort(nodebp→data, nodebp→size, sizeof(nodebp→data[0]),
cmpData); // Sort the data in time
   Tree = SplitBP(nodebp, Tree); // Split the node
    return Tree;
}
// Split the node when the node is full
```

```
// nodebp: the appropriate node where the term will be inserted
// Tree: B+ tree containing the inverted index
BplusTree SplitBP(NodeBP nodebp, BplusTree Tree) {
   if (!nodebp→childrenSize && nodebp→size ≤ ORDER // If
the node is not full
       | nodebp→childrenSize && nodebp→size < ORDER) {
(consider both leaf node and non-leaf node),
                                                          // do
       return Tree;
nothing!
   }
    // lnodebp, rnodebp: the left and right part of the split
node
    // tmpNodebp: store the node temporarily
    // parent: the parent node of nodebp
   NodeBP lnodebp, rnodebp, tmpNodebp, parent;
   int cut; // The position of the middle data
   int i, j;
   parent = nodebp→parent;
    if (!parent) { // If the node has no parent(i.e. this node
is the root),
         tmpNodebp = CreateBP(); // create a new node as the
parent(and also the root of the tree)
       parent = (NodeBP)malloc(sizeof(struct nodebp));
       Tree = parent = tmpNodebp;
   }
   lnodebp = CreateBP();
    rnodebp = CreateBP();
    lnodebp→parent = rnodebp→parent = parent; // Connect the
two parts with the parent node
    if (!nodebp→childrenSize) { // If the node is the leaf
node
       cut = LEAFCUT;
       for (i = 0; i < cut; i++) { // Assign the data in the
```

```
left part of original node to lnodebp
             lnodebp→data[i] = nodebp→data[i];
        lnodebp→size = cut;
         for (j = cut; j < nodebp \rightarrow size; j ++) \{ // Assign the
data in the right part of original node to rnodebp
             rnodebp \rightarrow data[j - cut] = nodebp \rightarrow data[j];
        rnodebp→size = nodebp→size - cut;
    } else {
                                      // If the node is the non-leaf
node
        cut = NONLEAFCUT;
         for (i = 0; i \le cut; i++) { // Assign the data and
children in the left part of original node to lnodebp
             if (i \neq cut)
                 lnodebp \rightarrow data[i] = nodebp \rightarrow data[i];
             lnodebp→children[i] = nodebp→children[i];
             lnodebp→children[i]→parent = lnodebp;
        }
        lnodebp→size = cut;
        lnodebp→childrenSize = cut + 1;
          // Assign the data and children in the right part of
original node to rnodebp
        for (j = cut + 1; j < nodebp \rightarrow size; j \leftrightarrow) {
             rnodebp \rightarrow data[j - cut - 1] = nodebp \rightarrow data[j];
        for (j = cut + 1; j < nodebp \rightarrow childrenSize; j \leftrightarrow) {
             rnodebp→children[j - cut - 1] = nodebp→children[j];
             rnodebp→children[j - cut - 1]→parent = rnodebp;
        }
        rnodebp→size = nodebp→size - cut - 1;
        rnodebp→childrenSize = nodebp→childrenSize - cut - 1;
    }
    // Assign the middle data in the original node to its parent
```

```
parent→data[parent→size++] = nodebp→data[cut];
   if (parent→childrenSize) { // If the parent has children(not
be created newly)
       for (i = 0; i < parent \rightarrow childrenSize; i++) {
            if (parent→children[i] = nodebp) { // Replace
the original node with lnodebp
               parent→children[i] = lnodebp;
               break;
           }
       }
   } else { // newly created parent
        parent→children[parent→childrenSize++] = lnodebp;
Insert the lnodebp
    parent→children[parent→childrenSize++] = rnodebp;
                                                             //
Insert the rnodebp
    // Sort the data and children of the parent
     gsort(parent→data, parent→size, sizeof(parent→data[0]),
cmpData);
   qsort(parent→children, parent→childrenSize, sizeof(parent-
>children[0]), cmpNodeBP);
   free(nodebp); // Free the memory of the original node
   Tree = SplitBP(parent, Tree); // Continue spliting the upper
node
   return Tree;
}
// Print the B+ tree(level-order traversal)
// T: B+ tree containing the inverted index
void PrintBPTree(BplusTree T) {
   int i;
   NodeBP nodebp; // The node obtained from the queue
   QueueBP q; // The queue containing the nodes from
B+ tree
```

```
printf("B+ Tree of Inverted Index:\n");
   q = CreateQueueBP(); // Create an empty queue
   EnqueueBP(T, q); // Put the root of the tree into the
queue first
   EnqueueBP(NULL, q); // Put the NULL pointer, for creation
of newline
   while (q \rightarrow size) { // If the queue isn't empty, repeat the
following steps
       nodebp = DequeueBP(q); // Get the front node
       time to add a newline
           printf("\n");
            if (q \rightarrow size) { // If the queue isn't empty,
continue add a new NULL pointer
              EnqueueBP(NULL, q);
       } else {
          printf("[");  // Print the node's data(just the
term)
          for (i = 0; i < nodebp \rightarrow size; i \leftrightarrow) {
              if (!i) {
                  printf("%s", nodebp→data[i]→term);
              } else {
                  printf(", %s", nodebp→data[i]→term);
              }
          printf("]");
       }
      then put its children into the queue
          for (i = 0; i < nodebp \rightarrow childrenSize; i++) {
              EnqueueBP(nodebp→children[i], q);
           }
       }
   }
}
```

```
// Create the queue
QueueBP CreateQueueBP() {
    QueueBP Q = (QueueBP)malloc(sizeof(struct queuebp));
    Q \rightarrow size = 0;
    Q \rightarrow front = Q \rightarrow rear = 0;
    return Q;
}
// Put the node of B+ tree into the queue
// nodebp: the newly added node
// Q: the queue
void EnqueueBP(NodeBP nodebp, QueueBP Q) {
     if (Q→size ≥ SIZE) { // If the queue is full, enqueue
operation fails
        printf("Full B+-tree-item queue!\n");
        exit(1);
    }
    Q \rightarrow data[Q \rightarrow rear ++] = nodebp; // Add new node
    Q→size++;
}
// Get the front node and delete it from the queue
// 0: the queue
NodeBP DequeueBP(QueueBP Q) {
    if (!Q→size) { // If the queue is empty, dequeue
operation fails
        printf("Empty B+-tree-item queue!\n");
        exit(1);
    }
    NodeBP returnNodeBP = Q \rightarrow data[Q \rightarrow front ++]; // Get the front
node
    Q→size--;
                    // Delete the node from queue
    return returnNodeBP;
}
// Create the poslist
PosList CreatePL() {
```

```
PosList L;
    L = (PosList)malloc(sizeof(struct poslist));
    L \rightarrow size = 0;
    L→front = (PosData)malloc(sizeof(struct posdata));
    L→rear = (PosData)malloc(sizeof(struct posdata));
    L \rightarrow front = L \rightarrow rear;
    L \rightarrow rear \rightarrow pos = -1; // Distinguish from other nodes
    return L;
}
// Add new position
// pos: the position
// L: the position list
void EnqueuePL(int pos, PosList L) {
    if (L \rightarrow rear \rightarrow pos \neq pos) { // If it's a new position
         PosData tmp = (PosData)malloc(sizeof(struct posdata));
         if (!tmp) {
              printf("Fail to create a new position data!\n");
              exit(1);
         } // Insert the new one in the position list
         tmp \rightarrow pos = pos;
         tmp \rightarrow time = 1;
         tmp \rightarrow next = L \rightarrow rear \rightarrow next;
         L \rightarrow rear \rightarrow next = tmp;
         L \rightarrow rear = tmp;
         L→size++;
    } else { // Otherwise, just increment the frequency
         L→rear→time++;
    }
}
// Retrieve all position in the list
// L: the position list
int ** RetrievePL(PosList L) {
      if (!L→size) { // If the list is empty, retrieve
operation fails
```

```
printf("Empty position-data queue!\n");
        exit(1);
    }
    int i = 0, j;
    int ** posArr = (int **)malloc(sizeof(int *) * L→size);
   for (j = 0; j < L \rightarrow size; j \leftrightarrow) \{ // Memory Allocation for 2D
array
        posArr[j] = (int *)malloc(sizeof(int) * 2);
    }
    PosData cur = L→front→next;
    while (cur \neq NULL) { // Make a traversal in the position
list
       posArr[i][0] = cur→pos; // Get the specific info of the
position
        posArr[i][1] = cur→time;
        cur = cur→next;
        i++;
    }
    return posArr;
}
// Build a hash table
HashTb GenerateHashTb() {
    int i;
                                   // Mark the start and the end
    int pre, cur;
of one word
    HashTb H;
                                   // The hash table containing
the stop words
                                  // File pointer
   FILE * fp;
   char fname[MAXWORDLEN];  // File name
    char tmp[MAXREADSTRLEN]; // Memory space storing the
reading data temporarily
                                  // Term(or word)
    char * term;
   H = InitHashTb();
                                   // Initialization
```

```
strcpy(fname, STOPWORDPATH); // Use default
path(stop_words.txt)
   fp = fopen(fname, "r");  // Open the file
   if (!fp) {
       printf("Fail to open the file of stopwords!\n");
       exit(1);
   }
    while (fgets(tmp, MAXREADSTRLEN - 1, fp) ≠ NULL) { //
Continue reading the file, until arrive at the end of file
       pre = cur = 0;
                                              // Initialization
       all characters in the tmp string
           if (!isalpha(tmp[i])) {
                                              // Maybe it's
time to record a word
              cur = i:
              if (cur > pre) {
                                              // Legitimate
situation
                   term = (char *)malloc(sizeof(char) * (cur -
pre + 1));
                  strncpy(term, tmp + pre, cur - pre);
                  term[cur - pre] = '\0';
                  InsertHashSW(term, H);
                                          // Insert the
new term
              }
              pre = cur + 1;
           }
       }
       // Handle the last possible word in the tmp string
       if (!cur || pre > cur && pre \neq i) {
           cur = i;
            term = (char *)malloc(sizeof(char) * (cur - pre +
1));
           strncpy(term, tmp + pre, cur - pre);
           term[cur - pre] = '\0';
           InsertHashSW(term, H);
                                               // Insert the
```

```
new term
        }
    }
    fclose(fp);
    return H;
}
// Initialization of the hash table
HashTb InitHashTb() {
    HashTb H;
                         // Hash table
    int i;
      H = (HashTb)malloc(sizeof(struct hashtb)); // Memory
allocation for the whole table
    if (H = NULL) {
        printf("Fail to create a hash table for stopwords!\n");
        exit(1):
    }
    H→size = STOPWORDSUM; // maxixum size
    for (i = 0; i < H \rightarrow size; i \leftrightarrow) {
        H→data[i] = (HashSW)malloc(sizeof(hashsw));
         if (H→data[i] = NULL) { // Memory allocation for
cells
             printf("Fail to create a hash table for stopwords!
\n");
           exit(1);
        }
           H→data[i]→stopword = (string)malloc(sizeof(char) *
MAXWORDLEN);
        H \rightarrow data[i] \rightarrow info = Empty;
    }
    return H;
}
// Find the stopwords or other words in the hash table
// stopword: stop word
```

```
// H: hash table containing the stop words
// justSearch: find the term without subsequent insertion
int FindHashSW(string stopword, HashTb H, bool justSearch) {
    int pos;
                                              // Appropraite position
    int collisionNum = 0;
                                                 // collision number,
for quadratic probe
    pos = HashFunc(stopword, H→size); // Use hashing function
first
    // Collision occurs!
     while (H \rightarrow data[pos] \rightarrow info \neq Empty \&\& strcmp(H \rightarrow data[pos] -
>stopword, stopword)) {
         pos += 2 * ++collisionNum - 1; // Quadratic probe
         if (pos \geq H\rightarrowsize)
             pos -= H→size;
         if (justSearch && H \rightarrow data[pos] \rightarrow info = Empty) {
             return -1:
         }
    }
    return pos;
}
// Insert a new stopword in hash table
// stopword: stop word
// H: hash table containing the stop words
void InsertHashSW(string stopword, HashTb H) {
    int pos;
    pos = FindHashSW(stopword, H, false); // Find the correct
position
     if (H \rightarrow data[pos] \rightarrow info \neq Legitimate) // Insert the stop
word
    {
         H \rightarrow data[pos] \rightarrow info = Legitimate;
         strcpy(H→data[pos]→stopword, stopword);
    }
}
// Hashing function
// stopword: stop word
```

```
// size: the maximum size of the hash table
int HashFunc(string stopword, int size) {
    unsigned int val = 0;
    while (*stopword \neq '\0')
         val = (val << 5) + *stopword++; // Generate the hash</pre>
value from every character in the string
    return val % size;
}
// Print hash table
void PrintHashTb(HashTb H) {
    int i;
    printf("Stopwords in hash table:\n");
    for (i = 0; i < H \rightarrow size; i \leftrightarrow) {
        if (H \rightarrow data[i] \rightarrow info \neq Empty) {
             printf("%d: %s\n", i, H→data[i]→stopword);
        }
    }
    printf("\n");
}
// Comparison functions used in qsort()
int cmpData(const void * a, const void * b) {
    const Data dataA = *(const Data*)a;
    const Data dataB = *(const Data*)b;
    return strcmp(dataA→term, dataB→term);
}
int cmpNodeBP(const void * a, const void * b) {
    const NodeBP nodebpA = *(const NodeBP*)a;
    const NodeBP nodebpB = *(const NodeBP*)b;
       return strcmp(nodebpA\rightarrowdata[0]\rightarrowterm, nodebpB\rightarrowdata[0]-
>term);
}
// wstring ←→ char *, for word stemming
```

```
std::wstring chararrToWstring(char * st) {
    std::string tmp(st);
    std::wstring_convert<std::codecvt_utf8<wchar_t>> converter;
    std::wstring wstr = converter.from_bytes(tmp);
   return wstr;
}
char * wstringToChararr(std::wstring wst) {
   std::wstring_convert<std::codecvt_utf8<wchar_t>> converter;
   std::string tmp = converter.to_bytes(wst);
   char * st = new char[tmp.size() + 1];
    strcpy(st, tmp.c_str());
   return st;
}
// Word Stmming wrapper
string WordStem(string term) {
   std::wstring term_wstr;
                                          // the wstring form of
the term
      stemming::english_stem<> StemEnglish; // Word stemming
function(a little clumsy)
    term_wstr = chararrToWstring(term);
               transform(term_wstr.begin(), term_wstr.end(),
term_wstr.begin(), ::tolower);
   StemEnglish(term_wstr);
   term = wstringToChararr(term_wstr);
   return term;
}
// Print the ticks and duration, for -tr or --time function
void PrintTime(clock_t start, clock_t end) {
                    // ticks
   clock_t tick;
   double duration; // duration(unit: seconds)
   int iterations;
```

```
iterations = ITERATIONS;
tick = end - start;
duration = ((double)(tick)) / CLOCKS_PER_SEC;
printf("Iterations: %d\n", iterations);
printf("Ticks: %lu\n", (long)tick); // Print the info
printf("Duration: %.2fs\n", duration);
}
```

## 5.5 invIndexTest.cpp

```
#include "invIndexHeader.h"
#include "wordStem/english_stem.h"
#include <stdbool.h>
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include <string>
#include <time.h>
parameters
   int i;
   bool isTest = false;
                                        // Whether open test
mode(-t or --test)
   bool Print = false;
                                       // Whether open print
mode(-p or --print)
   bool isFound;
                                      // Whether the term is
found in the inverted index
   bool containStopWords = false;  // Whether open stopword
mode(-s or --stopwords)
   bool timeRecord = false;
                                      // Whether open time
record mode(-tr or --time)
   int findCnt = 0;
                                       // The time of finding
a single word for find mode(-f=n or --find=n)
   char * pos;
                                       // Get the number after
`=` in parameters
    char tmp[MAXWORDLEN];
                                          // Store the input
string temporarily
    char * word;
                                           // The word to be
```

```
searched
                                        // Duration of running
   double duration;
a function
    std::wstring word_wstr;
                                           // wstring form of
      stemming::english_stem<> StemEnglish; // Word stemming
function
                                       // Inverted Index
   BplusTree InvIndex;
   clock_t start, end, tick;
                                        // Record the start
and the end of the clock
   for (i = 1; i < argc; i++) { // Read the parameters
       if (!strcmp(argv[i], "--test") || !strcmp(arqv[i], "-t"))
{ // Test mode
           isTest = true;
        } else if (strstr(argv[i], "--find") || strstr(argv[i],
"-f")) { // Find mode
           if ((pos = strchr(arqv[i], '='))) {
                 findCnt = atoi(pos + 1); // Get the number
behind `=`
               if (!findCnt) {
                   printf("Wrong Number!\n");
                   exit(1);
               }
           } else {
               findCnt = 1; // Use default number
      } else if (!strcmp(argv[i], "--print") || !strcmp(argv[i],
"-p")) { // Print mode
           Print = true:
            } else if (!strcmp(arqv[i], "--stopwords") || !
strcmp(arqv[i], "-s")) { // Stopword mode
           containStopWords = true;
       } else if (!strcmp(argv[i], "--time") || !strcmp(argv[i],
"-tr")) { // Time record mode
           timeRecord = true:
       } else { // Error
           printf("Wrong Parameter!\n");
           exit(1);
```

```
}
    }
    if (!timeRecord) { // No time record
        InvIndex = InvertedIndex(isTest, containStopWords);
    } else { // Time record
        char dir[MAXREADSTRLEN];
        strcpy(dir, IISPEEDTESTDIR);
        start = clock();
        for (i = 0; i < ITERATIONS; i++) {
            InvIndex = CreateBP();
          InvIndex = fileTraversaler(InvIndex, dir, NULL, isTest,
containStopWords);
        }
        end = clock();
        PrintTime(start, end);
    }
    if (Print) { // Print the B+ tree
        PrintBPTree(InvIndex);
    }
   if (InvIndex→size && findCnt) { // Search the single word
        word = (char *)malloc(sizeof(char) * MAXWORDLEN);
         printf("\nFinding Words Mode(only supports single word
finding):\n");
        for (i = 0; i < findCnt; i++) { // For every search}
            isFound = false;
            printf("Find %d: ", i + 1);
            scanf("%s", tmp);
            strcpy(word, tmp);
            // Word stemming
            word = WordStem(word);
            // Find the word
            if (!timeRecord) { // No time record
                FindBP(word, -1, InvIndex, &isFound, findCnt);
            } else { // Time record
```

```
start = clock();
    FindBP(word, -1, InvIndex, &isFound, findCnt);
    end = clock();
    PrintTime(start, end);
}

// If not found, then give relevant information
    if (!isFound) {
        printf("Sorry, no such word in the inverted index!

\n");
    printf("-----\n");
    }
}

return 0;
}
```

## 5.6 invIndexSearch.cpp

```
/************
* The following code is the implementation of the inverted index
search algorithm.
* The algorithm takes a query from the user and searches for the
documents that contain all the valid query words.
* The algorithm uses the B+ tree to search for the documents and
the inverted index to get the word frequencies.
* The algorithm also uses the stop list to ignore the common
words and the stemming algorithm to reduce the words to their
root form.
**************
#include <iostream>
                                  // for input/output
#include <fstream>
                                  // for file input/output
#include <string>
#include <string.h>
#include <map>
                                  // for map data structure
#include <unordered_map>
                                  // for unordered_map data
structure (hash)
```

```
#include <algorithm>
                                   // for sorting
#include <cctype>
#include <cmath>
                                   // for log()
#include "wordStem/english_stem.h" // for stemming
#include "invIndexHeader.h"
#define DOCTOTALNUM 992
                                 // total number of documents
//hash table: {word:wordIdf}
std::unordered_map<std::wstring, double> wordIdf;
//hash table: {docId:fileWordCount}
std::unordered_map<std::wstring, int> fileWordsNum;
//hash table: {word:bool} PS: Bool is just a placeholder and has
no actual function.
std::unordered_map<std::wstring, bool> stopWords;
stemming::english_stem<> StemEnglish;
/***********
* This function is the main function for searching.
* In this function, we analyzes the guery
* and search for the documents that contain all the valid query
words.
* Input: T: Inverted index (B+ tree)
         pageSize: number of documents to be displayed per query
         threshold: the proportion of search terms to the total
number of terms
**************
void search(std::string query, BplusTree T, int pageSize, double
threshold)
{
   /*load the query and analyze it*/
    //vector: Elements are {word:wordIdf}
   std::vector<std::pair<std::wstring,double>> queryWord;
   std::wstring word=L"";
    //Iterate through each character in the string and extract
the words
   for(char &c : query)
   ₹
        // if the character is alpha or number, add it to the
```

```
word. It solve the problem of the Punctuation and Whitespace
        if(isalnum(c))
            word += tolower(c);
        else if (word.length() > 0)
        ₹
            std::wstring tmp = word; // save original word
            StemEnglish(word);
            // check if the word is in the stop list
            auto it = stopWords.find(word);
            if(it \neq stopWords.end())
               // if the word is in the stop list, print a warning
message and ignore it.
              std::wcout << L"\033[1;31mWarning: \033[0m\"" << tmp
<< L"\" is in the stop list and will be ignored." << std::endl;</pre>
            else
            {
                // if the word is not in the stop list, check if
it is in the inverted index.
                auto it2 = wordIdf.find(word);
                if(it2 = wordIdf.end())
                    // if the word is not in the inverted index,
print a warning message and ignore it.
                   std::wcout << L"\033[1;31mWarning: \033[0m\""
<< tmp << L"\" is not in the inverted index and will be ignored."</pre>
<< std::endl;</pre>
                else
                     // if the word is in the inverted index, add
it to the queryWord vector.
                    queryWord.push_back({word,wordIdf[word]});
            word = L""; // reset the word
        }
    }
    // Sort the queryWord vector by wordIdf in descending order
                         sort(queryWord.begin(),queryWord.end(),
[](const
              std::pair<std::wstring,double>&
                                                    a,
std::pair<std::wstring,double>& b){return a.second > b.second;});
    /* Search for documents that contain all the query words */
```

```
int cntForDoc = 0;
                                                                //
counter for pageSize
    int cntForWord = (int)(queryWord.size() * threshold);
                                                               //
counter for threshold
    //control the size of cntForWord, to avoid bugs
    if (cntForWord = 0)
        cntForWord = 1;
    else if (cntForWord > queryWord.size())
        cntForWord = queryWord.size();
    char * wordForSearch;
                                                               //
char array for search
    //if the query is empty or only has stop words, return
    if(queryWord.size() = 0)
    {
       std::cout << "\033[1;31m0ps, your query is in StopList or</pre>
empty, so there are no documents retrieved. \033[0m" << std::endl;
        return;
    }
    //if the guery has only one valid word, search for it in the
inverted index
    //here, we only use "Tf" to sort the documents in rusult
    else if(queryWord.size() = 1)
    {
        //vector: Elements are {docId:tf}
        std::vector<std::pair<int,double>> posVec;
        std::cout << "Your query has only one valid word, so we</pre>
will search for it in the inverted index." << std::endl;
          std::cout << "The word was found in files below:"<</pre>
std::endl:
        printf("\033[1,32mFile name
                                                    \tTf\033[0m\n"]);
         //don't need to traverse here, but I wrote the case of
multiple words first, so I reused it directly
        for(auto &p : queryWord)
        {
            wordForSearch = wstringToChararr(p.first);
            //find all the positions of the word in the B+ tree
            posVec = FindBP2(wordForSearch, -1, T);
            //sort the positions by tf in descending order
```

```
sort(posVec.begin(),posVec.end(),[](const
std::pair<int,double>& a, const std::pair<int,double>& b){return
a.second > b.second;});
            //print the file name and tf of the word
            for(auto &pos : posVec)
            {
                if(cntForDoc = pageSize)
                    break;
                printf("\033[34m%-25s\t%lf\033[0m\n", docNames[pos.first],
                cntForDoc++;
            }
        }
    // if the query has multiple valid words, search for them in
the inverted index
    // Here, we use "Tf-Idf" to sort the documents in result
    else
    {
        std::cout << "Your query has multiple valid words, so we</pre>
will search for them in the inverted index." << std::endl;
         std::cout << "The words were found in files below:"<</pre>
std::endl;
        //hash table: {docId : sum of tf-idf}
        std::unordered_map<int, double> freqMap;
        //traverse the queryWord vector and search for each word
in the inverted index
        bool flag=false;
        for(auto &p : queryWord)
        {
            if(cntForWord = 0)
                break;
            wordForSearch = wstringToChararr(p.first);
            auto currentPosVec = FindBP2(wordForSearch, -1, T);
             //Transfer the values in the vector to a temporary
hash table, mainly for efficiency reasons.
            //The temporary hash table only contains the pair of
docId and idf.
```

```
std::unordered_map<int, double> currentDocIdMap;
            if(flag = false)
            {
                for(auto &pos : currentPosVec)
                 freqMap[pos.first] += pos.second * p.second;
tf-idf = tf * idf
                flag = true;
            }
            else
            {
                for(auto &pos : currentPosVec)
                      currentDocIdMap[pos.first] = pos.second *
p.second;
                if(freqMap.size()>0)
                        //remove "it++" in "for(;;)"" to remove
duplicates
                         for (auto it = freqMap.begin(); it ≠
freqMap.end(); )
                    {
                          if (currentDocIdMap.find(it→first) =
currentDocIdMap.end())
                            it = freqMap.erase(it);//safe deletion
                        else
                        {
                               it→second += currentDocIdMap[it-
>first];
                            #it;
                        }
                    }
                }
                else
                    break;
            cntForWord--;
        }
         //if there is no document that contains all the query
words, return
        if(freqMap.size() = 0)
```

```
{
            std::cout << "Ops, your query is not in the inverted</pre>
index, so there are no documents retrieved. " << std::endl;
           std::cout << "Please try again with different query."</pre>
<< std::endl;</pre>
            return;
        //if there is at least one document that contains all the
query words, sort them by tf-idf in descending order
        else
        {
             printf("\033[1;32mFile name
                                                             \tTf-
Idf\033[0m\n");
             //convert the unordered_map to a vector and sort it
by tf-idf in descending order
                              std::vector<std::pair<int,double>>
sortedFreqVec(freqMap.begin(), freqMap.end());
           std::sort(sortedFreqVec.begin(), sortedFreqVec.end(),
[](const std::pair<int, double>& a, const std::pair<int, double>&
b) {return a.second > b.second; });
            //print the file name and tf-idf of the word
            for(auto &p : sortedFreqVec)
            {
                if(cntForDoc = pageSize)
                printf("\033[34m%-25s\t%lf\033[0m\n",docNames[p.first],p.
                cntForDoc++;
            }
        }
    }
}
// similar to FindBP() in invIndexFunc.cpp, but desgined just for
searching
// return <word, tf>
std::vector<std::pair<int,double>> FindBP2(string
                                                     term,
                                                               int
docCnt, BplusTree T)
{
```

```
int i;
    // if the tree is empty, return an empty vector
    if (!T)
        return std::vector<std::pair<int,double>>();
    else if (!T→childrenSize)
        return isSameTerm2(term, docCnt, T);
    int pos = -1;
    // find the position of the term in the current node
    for (i = 0; i < T \rightarrow size; i \leftrightarrow) {
        if (strcmp(term, T \rightarrow data[i] \rightarrow term) < 0) {
             pos = i;
             break;
        }
    }
    if (pos = -1) {
        pos = i;
    }
    return FindBP2(term, docCnt, T→children[pos]);
}
//similar to isSameTerm() in invIndexFunc.cpp, but designed just
for searching
//return <docId, tf>
std::vector<std::pair<int,double>> isSameTerm2(string term, int
docCnt, NodeBP nodebp)
{
    int i;
    // vector: Elements are {docId:idf}
    std::vector<std::pair<int,double>> posVec;
    if (nodebp→size)
    {
        for (i = 0; i < nodebp \rightarrow size; i++) {
             if (!strcmp(term, nodebp→data[i]→term))
             {
```

```
PosList poslist = nodebp→data[i]→poslist;
                posVec = RetrievePL2(poslist);
                break;
            }
        }
    }
    return posVec;
}
//similar to RetrievePL() in invIndexFunc.cpp, but designed just
for searching
//return <docId, tf>
std::vector<std::pair<int,double>> RetrievePL2(PosList L)
{
    // if the position-data queue is empty, return an empty vector
    if (!L\rightarrowsize) {
        printf("Empty position-data queue!\n");
        exit(1);
    }
    double tf;
    std::vector<std::pair<int,double>> posVec;
    PosData cur = L→front→next;
    while (cur ≠ NULL) {
          std::wstring docName = chararrToWstring(docNames[cur-
>pos]);
        // calculate tf : tf = word freq/total words in file
        tf = (double)cur→time / (double)(fileWordsNum[docName]);
        posVec.push_back({cur→pos, tf});
        cur = cur→next;
    }
    return posVec;
}
/*The functions below load the wordIdf, fileWordsNum, and
stopWords from files*/
```

```
void loadWordIdf(std::string filePath)
{
    std::wifstream infile;
    std::wstring word;
    int freq;
    double idf;
    infile.open(filePath,std::ios::in);
    //if the file is empty, return
    if(!infile.is_open())
    {
        std::cout << "Error: unable to open file " << filePath <<
std::endl;
        return;
    //read the word and its frequency from the file and calculate
its idf
    while(infile >> word >> freq)
    {
        //calculate idf : idf = log(total number of documents/(1
+ number of documents contain the word))
        idf = log((double)DOCTOTALNUM/(double)(1+freq));
        wordIdf[word] = idf;
    }
    infile.close();
    std::cout << "Word Idf loaded successfully." << std::endl;</pre>
}
void loadFileWordsNum(std::string filePath)
{
    std::wifstream infile;
    std::wstring filename;
    int num;
    infile.open(filePath,std::ios::in);
    //if the file is empty, return
    if(!infile.is_open())
    {
        std::cout << "Error: unable to open file " << filePath <<
std::endl;
        return;
```

```
}
    //read the file name and its word count from the infile
    while(infile >> filename >> num)
    {
        filename += L".txt";
        fileWordsNum[filename] = num;
    }
    infile.close();
   std::cout << "The word count of every file loaded successfully."</pre>
<< std::endl;</pre>
}
void loadStopWords(std::string filePath)
{
    std::wifstream infile;
    std::wstring word;
    infile.open(filePath,std::ios::in);
    //if the file is empty, return
    if(!infile.is_open())
        std::cout << "Error: unable to open file " << filePath <</pre>
std::endl;
        return;
    }
    while(infile >> word)
        stopWords[word] = true;
    std::cout << "Stop words loaded successfully." << std::endl;</pre>
}
```

# 5.7 search\_test.cpp

```
int main()
{
    clock_t start, end;
    //load inverted index and other necessary resources
    std::cout << "Please wait for resources to load and inverted</pre>
index to be established..."<<std::endl;</pre>
     loadWordIdf(WORDDOCSPATH);
                                                 //load word-idf
dictionary
   loadFileWordsNum(FILEWORDCOUNTPATH); //load file-word count
dictionary
    loadStopWords(STOPWORDPATH);
                                              //load stop words
list
    BplusTree InvIndex = CreateBP();  //create B+ tree for
inverted index
    InvIndex = fileTraversaler(InvIndex, (char *)SHAKESPEAREDIR,
(char *)"", false, true);
        std::cout << "Inverted Index has been established</pre>
successfully!"<<std::endl;</pre>
                                          std::cout
                                                               <<
std::endl;
    std::cout << "Welcome to ShakespeareFinder's test program!"</pre>
<< std::endl;</pre>
   std::cout << "The program tests the performance of the search</pre>
engine in different situations" << std::endl;</pre>
    //test loop begin
   testbegin:
   std::cout << "Please enter the serial number of the input you</pre>
want to test: " << std::endl;</pre>
    int serial_num;  //serial number of input file
    std::cin >> serial_num;
             std::string input_file_path = root_path +
std::to_string(serial_num); //input file path
    std::ifstream input_file;
                                                          //input
file object
    input_file.open(input_file_path, std::ios::in); //open
```

```
input file
    // if input file is not found, return
    if (!input_file.is_open())
        std::cout << "Invalid input serial number!" << std::endl;</pre>
    // if input file is found, continue
    else
    {
        //read input parameters
        int pageSize, k;
        double threshold;
        std::cout << "Input file opened successfully!" << std::endl;</pre>
         std::cout << "Please enter the pageSize, threshold and</pre>
the times of cycles : " << std::endl;
        std::cin >> pageSize >> threshold >> k;
        std::string query;
        std::getline(input_file, query);
        query += " ";
        //test performance
        start = clock();
        for(int i=0; i<k; i++)
        {
               std::cout << "Cycle " << i+1 << " starts..." <<
std::endl; //print cycle number
            search(query, InvIndex , pageSize, threshold);
        }
        end = clock();
        PrintTime(start,end);
    }
    //ask user if they want to test another input
    std::cout << "Do you want to test another input? (y/n)" <<
std::endl;
    char choice;
    std::cin >> choice;
    if(choice = 'y')
    {
        getchar();
```

```
goto testbegin; //jump to testbegin
}
else
  return 0;
}
```

#### 5.8 search\_main.cpp

```
/*************
* This is the main function of the search engine.
* It first loads the inverted index and the necessary resources.
* Then it enters a loop to repeatedly search for a query until
the user chooses to exit.
*****************
#include <iostream>
#include "invIndexHeader.h"
int main()
₹
    //load necessary resources and inverted index
    std::cout << "Please wait for resources to load and inverted</pre>
index to be established..."<<std::endl;</pre>
    loadWordIdf(WORDDOCSPATH);
                                               //load word-idf
dictionary
   loadFileWordsNum(FILEWORDCOUNTPATH); //load file-word count
dictionary
    loadStopWords(STOPWORDPATH);
                                            //load stop words
list
    BplusTree InvIndex = CreateBP();  //create B+ tree for
inverted index
   InvIndex = InvertedIndex(); // fileTraversaler(InvIndex, (char
*)SHAKESPEAREDIR, (char *)"", false, true);
        std::cout << "Inverted Index has been established</pre>
successfully!"<<std::endl;</pre>
                                         std::cout
                                                            <<
std::endl;
```

```
//set values for pageSize and threshold from user input
                   std::cout << "\033[1;32mWelcome
                                                               to
***ShakespeareFinder***\033[0m" << std::endl;
   std::cout << "\033[34mPlease set the maximum number of records</pre>
to be displayed per query\033[0m" << std::endl;
    int pageSize = 10;
                                //default value
    std::cout << "Size(int): ";</pre>
    std::cin >> pageSize;
    std::cout << "\033[34mPlease set the threshold (0-1) of the
query:\033[0m" << std::endl;
    std::cout << "Threshold(double): ";</pre>
    double threshold = 0.8;  //default value
    std::cin >> threshold;
   qetchar();
                               //clear the buffer to avoid input
error
    //start the search loop
    Searchbegin:
    std::cout << "Please enter your query:" << std::endl;</pre>
    std::string query;
    /* Read query from user and analyse it*/
    std::getline(std::cin,query);
                                 // add a space at the end of
    query += " ";
the query to avoid errors
    search(query, InvIndex, pageSize, threshold);
    //ask user if they want to search again
     std::cout << "Do you want to search again?\033[1;31m(y/</pre>
n)\033[0m: ";
    char choice:
    std::cin >> choice;
    if(choice = 'y' || choice = 'Y')
    {
        getchar();
        goto Searchbegin;
    }
    //exit the program
    else
    {
```

```
std::cout << "\033[1;32mThank you for using
ShakespeareFinder!" << std::endl << "Have a nice day~\033[0m" <<
std::endl;
    return 0;
}
</pre>
```

# References

 $[1] \ Blake-Madden, \ Oleander Stemming Library, \ \underline{https://github.com/Blake-Madden/Oleander Stemming Library}$ 

#### Declaration

We hereby declare that all the work done in this project titled "Roll Your Own Mini Search Engine" is of our independent effort as a group.