

BLOOM LEVEL SETTING

report for the CSE204:Design and Analysis of Algorithm project

Sumbitted by

P.Bhavya sri(AP23110010929)

G.Dhanush(AP23110010867)

K.Vaishnavi(AP23110010888)

Bachelor of Technology

In

Computer Science and Engineering

School of Engineering and Sciences



Under the guidance of
Mr. PAMULA UDAY RAJU
Assistant Professor-Ad Hoc
Department of CSE
[November, 2024]

SRM University, AP, Neerukonda, Mangalagiri, Guntur.

Andhra Pradesh – 522 240

Department of Computer Science and Engineering
SRM University, AP



CERTIFICATE

*This is to certify that the Project report entitled “**Bloom level settings**” is being submitted by **Bhavya sri** (AP23110010929), **Dhanush** (AP23110010867) and **vaishnavi** (AP23110010888) students of Department of Computer Science and Engineering, SRM University, AP, in partial fulfillment of the requirement for Design and Analysis of Algorithms Lab for II-B. Tech (CSE), Semester III. carried out by him during the academic year 2024-2025.*

Signature of the Lab In charge

Mr. PAMULA UDAY RAJU

Assistant Professor-Ad Hoc

Department of CSE

Abstract

This project aims to implement an Outcome-Based Education (OBE) system at SRM-AP University, focusing on managing Bloom's Taxonomy levels through a user-friendly application. The system allows users to perform Create, Update, Retrieve, and Delete (CRUD) operations on Bloom entries, which include a Bloom Code, Level, and Description. The application employs efficient sorting and searching algorithms to organize and retrieve data effectively.

Two sorting algorithms are implemented: Quick Sort and Merge Sort, which are used to sort Bloom entries based on their codes. The system compares the performance of both algorithms, with Quick Sort providing an average time complexity of $O(n \log n)$ and a worst-case complexity of $O(n^2)$, while Merge Sort consistently performs at $O(n \log n)$. Additionally, two searching algorithms are utilized: Linear Search and Binary Search. Linear Search is used for unsorted data with a time complexity of $O(n)$, while Binary Search is applied after sorting, offering a faster time complexity of $O(\log n)$.

The application also features file handling capabilities, allowing data to be stored and retrieved from a text file ('bloom_setting.txt'), which ensures persistence across sessions. Through these modules, the system enables efficient data management and analysis, with the ability to display the time complexities of both sorting and searching algorithms.

The project demonstrates the use of various algorithms in practical scenarios within the educational context and highlights the benefits and trade-offs of each algorithm used in terms of time complexity and efficiency.

Table of Contents

○ List of diagrams	- 5
○ Introduction	- 6
○ Project Modules:	- 6
○ Architecture Diagram	- 6
○ Module Description	- 7
○ Programming Details	-7
○ Field/table details	-7
○ Algorithm Details	-8 to 18
○ (i)Sorting	
○ (ii)Searching	
○ (ii) Storing the details in a text file	
○ Source Code	
○ Comparison of Sorting Algorithms	-18
○ Comparison of Searching Algorithms	-19
○ Screen Shots	-20 to 22
○ Conclusion	-23

LIST OF DIAGRAMS

SNO	Name of diagram	page number
1	Architecture diagram	5
2	Table details	6
3	Screenshot outputs 1	18
4	Screenshot ouputs 2	19
5	Screenshot outputs 3	20

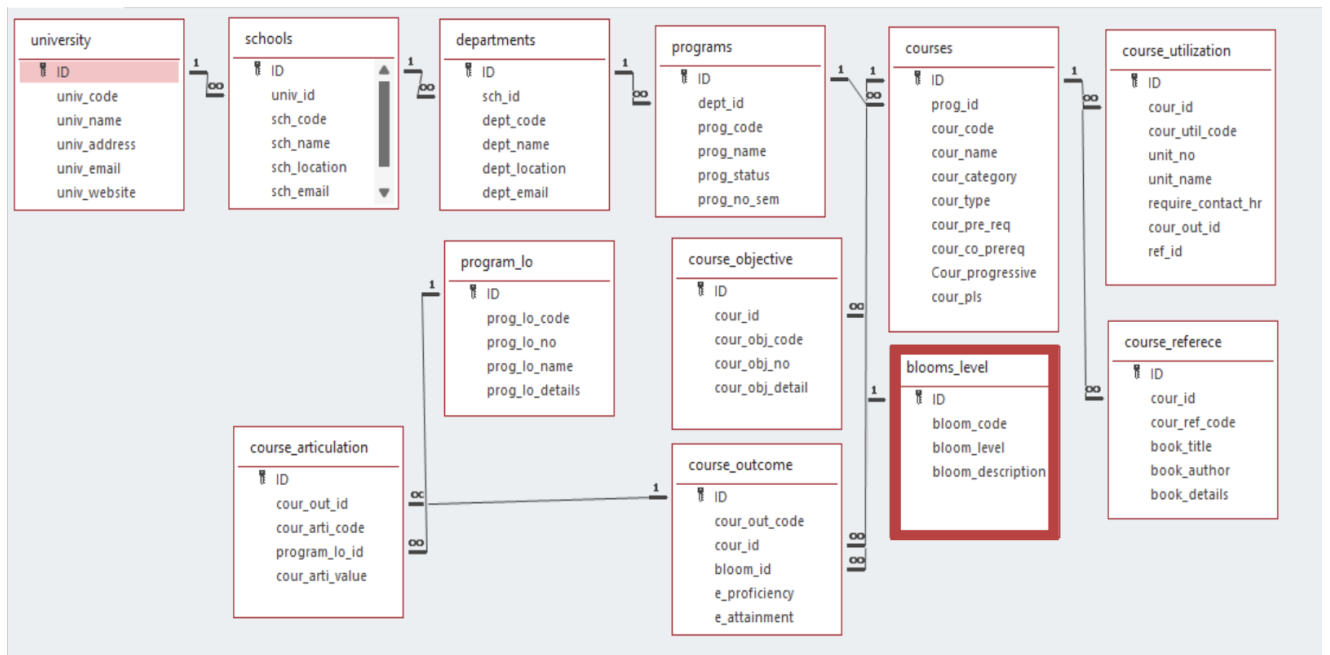
Introduction

Our University (herewith considered as SRM-AP) is going to implement OBE(Outcome Based Education) in their university and you assigned in the project to develop an application with any programming Language you are well versed and you were supposed to do searching and sorting using learned algorithms,comparing your sorting algorithm with any one of existing algorithm,displaying the time complexity of both algorithms and explaining advantages and disadvantages of the algorithm.

Project Modules:

1. Bloom Level setting
2. Search and Sorting Functions
3. Text File Handling

Architecture Diagram



Module Name: Bloom Level Setting

Module Description:

This module is used to create, Update, Retrieve, Delete (hereafter known as CRUD) details of the module and storing the details in the text file. You have to provide option for any of the operations of fields mentioned below according to algorithms given for you.

Programming Details naming conventions to be used:

File name: bloom_setting.txt

Function Naming Conventions:

- Create: titans_bloom_create
- Update: titans_bloom_update
- Retrieve: titans_bloom_retrieve
- Delete: titans_bloom_delete
- Sorting: titans_compare_sort_mergesort (Merge Sort) and titans_quick_sort (Quick Sort)
- Searching: titans_linear_search (Linear Search) and titans_compare_search_binarysearch (Binary Search)

- File Operations: titans_bloom_storing (for storing data)

Bloom Level Setting:Field/table details

Field name	Data type
id	int
bloom_code	String
bloom_level	String
bloom_des	String

Algorithm Details:

(i) Sorting

Sorting algorithms used:

- Quick Sort: Used for sorting Bloom entries by 'code'. Complexity: $O(n^2)$ worst case, $O(n \log n)$ average case.
- Merge Sort: Used for sorting Bloom entries by 'code'. Complexity: $O(n \log n)$.

(ii) Searching

Searching algorithms used:

- Linear Search: Simple search with $O(n)$ complexity.
- Binary Search: Searches after sorting, with $O(\log n)$ complexity.

Source Code (Full Implementation):

```
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
```



```

#define MAX 100

typedef struct {
    int id;
    char code[50];
    char level[100];
    char des[100];
} bloom;

bloom b[MAX];
int c = 0;
const char* FILE_NAME = "bloom_setting.txt";

void titans_load_from_file();
void titans_bloom_storing();
void titans_bloom_create();
void titans_bloom_update();
void titans_bloom_retrieve();
void titans_bloom_delete();
void titans_quick_sort(int low, int high);
int titans_partition(int low, int high);
void titans_compare_sort_mergesort(int left, int right);
void titans_merge(int left, int mid, int right);
int titans_linear_search(char *code);
int titans_compare_search_binarysearch(char *code);
void titans_bloom_search();
void titans_complexity_searching();
void titans_complexity_sorting();

void titans_load_from_file() {
    FILE *file = fopen(FILE_NAME, "r");
    if (file == NULL) {
        printf("File not found or unable to open: %s\n", FILE_NAME);
        return;
    }
    c = 0;
    while (fscanf(file, "%d %49s %99s %99[^\n]", &b[c].id, b[c].code, b[c].level, b[c].des) == 4) {
        c++;
    }
    fclose(file);
}

void titans_bloom_storing() {
    FILE *file = fopen(FILE_NAME, "w");
    if (file == NULL) {
        printf("Error opening file for writing: %s\n", FILE_NAME);
    }
}

```

```

        return;
    }
    for (int i = 0; i < c; i++) {
        fprintf(file, "%d %s %s %s\n", b[i].id, b[i].code, b[i].level, b[i].des);
    }
    fclose(file);
}

```

```

void titans_bloom_create() {
    if (c >= MAX) {
        printf("Bloom list is full!\n");
        return;
    }
    bloom bl;
    printf("Enter bloom ID: ");
    if (scanf("%d", &bl.id) != 1) {
        printf("Invalid input!\n");
        return;
    }
    printf("Enter bloom Code: ");
    scanf("%49s", bl.code);
    printf("Enter bloom Level: ");
    scanf(" %99[^\n]", bl.level);
    printf("Enter bloom Description: ");
    scanf(" %99[^\n]", bl.des);

    b[c++] = bl;
    titans_bloom_storing();
    printf("Bloom created successfully!\n");
}

```

```

void titans_bloom_update() {
    int id;
    printf("Enter bloom ID to update: ");
    if (scanf("%d", &id) != 1) {
        printf("Invalid input!\n");
        return;
    }

    for (int i = 0; i < c; i++) {
        if (b[i].id == id) {
            printf("Enter new bloom Code: ");
            scanf("%49s", b[i].code);
            printf("Enter new bloom Level: ");
            scanf(" %99[^\n]", b[i].level);
            printf("Enter new bloom Description: ");
            scanf(" %99[^\n]", b[i].des);

```

```

        titans_bloom_storing();
        printf("Bloom updated successfully!\n");
        return;
    }
}
printf("Bloom with ID %d not found.\n", id);
}

void titans_bloom_retrieve() {
    if (c == 0) {
        printf("No blooms available.\n");
        return;
    }
    printf("\nList of Blooms:\n");
    for (int i = 0; i < c; i++) {
        printf("ID: %d\nCode: %s\nLevel: %s\nDescription: %s\n\n",
            b[i].id, b[i].code, b[i].level, b[i].des);
    }
}

void titans_bloom_delete() {
    int id;
    printf("Enter bloom ID to delete: ");
    if (scanf("%d", &id) != 1) {
        printf("Invalid input!\n");
        return;
    }

    for (int i = 0; i < c; i++) {
        if (b[i].id == id) {
            for (int j = i; j < c - 1; j++) {
                b[j] = b[j + 1];
            }
            c--;
            titans_bloom_storing();
            printf("Bloom deleted successfully!\n");
            return;
        }
    }
    printf("Bloom with ID %d not found.\n", id);
}

int titans_partition(int low, int high) {
    char pivot[50];
    strcpy(pivot, b[high].code);
    int i = low - 1;

```

```

for (int j = low; j < high; j++) {
    if (strcmp(b[j].code, pivot) < 0) {
        i++;
        bloom temp = b[i];
        b[i] = b[j];
        b[j] = temp;
    }
}
bloom temp = b[i + 1];
b[i + 1] = b[high];
b[high] = temp;
return i + 1;
}

```

```

void titans_quick_sort(int low, int high) {
    if (low < high) {
        int pivot = titans_partition(low, high);
        titans_quick_sort(low, pivot - 1);
        titans_quick_sort(pivot + 1, high);
    }
}

```

```

void titans_merge(int left, int mid, int right) {
    int n1 = mid - left + 1;
    int n2 = right - mid;

    bloom L[n1], R[n2];

    for (int i = 0; i < n1; i++)
        L[i] = b[left + i];
    for (int i = 0; i < n2; i++)
        R[i] = b[mid + 1 + i];

    int i = 0, j = 0, k = left;
    while (i < n1 && j < n2) {
        if (strcmp(L[i].code, R[j].code) <= 0) {
            b[k] = L[i];
            i++;
        } else {
            b[k] = R[j];
            j++;
        }
        k++;
    }
}

```

```

while (i < n1) {
    b[k] = L[i];

```

```

        i++;
        k++;
    }

    while (j < n2) {
        b[k] = R[j];
        j++;
        k++;
    }
}

void titans_compare_sort_mergesort(int left, int right) {
    if (left < right) {
        int mid = left + (right - left) / 2;
        titans_compare_sort_mergesort(left, mid);
        titans_compare_sort_mergesort(mid + 1, right);
        titans_merge(left, mid, right);
    }
}

int titans_linear_search(char *code) {
    for (int i = 0; i < c; i++) {
        if (strcmp(b[i].code, code) == 0) {
            return i;
        }
    }
    return -1;
}

int titans_compare_search_binarysearch(char *code) {
    titans_compare_sort_mergesort(0, c - 1);
    int left = 0;
    int right = c - 1;

    while (left <= right) {
        int mid = left + (right - left) / 2;
        int cmp = strcmp(b[mid].code, code);
        if (cmp == 0) {
            return mid;
        } else if (cmp < 0) {
            left = mid + 1;
        } else {
            right = mid - 1;
        }
    }
    return -1;
}

```

```

void titans_complexity_searching()
{

    printf("Linear Search: time complexity is O(n).\n");

    printf("Binary Search: time complexity is O(log n).\n");

}

void titans_complexity_sorting()
{
    printf("Quick Sort: time complexity is O(n^2) for worst case[when last element is chosen as
pivot as the elements are already in sorted order] and O(n log n) for the average case.\n");

    printf("Merge Sort: time complexity is O(n log n).\n");

}

int main() {
    titans_load_from_file();

    int choice;
    while (1) {
        printf("\n1. Create Bloom\n2. Update Bloom\n3. Retrieve Blooms\n4. Delete Bloom\n5.
Compare Search Algorithm(binary search)\n6. Compare Sort
Algorithm(merge sort)\n7. Sort by Code (Quick Sort)\n8. Search by Code (Linear Search)\n9.
Complexity search\n10. Complexity Sort\n11. Exit\n");
        printf("Enter your choice: ");
        if (scanf("%d", &choice) != 1) {
            printf("Invalid input!\n");
            while (getchar() != '\n');
            continue;
        }

        switch (choice) {
            case 1:
                titans_bloom_create();
                break;
            case 2:
                titans_bloom_update();
                break;
            case 3:
                titans_bloom_retrieve();
                break;

```

```

case 4:
    titans_bloom_delete();
    break;
case 5:
{
    char code[50];
    printf("Enter bloom code to search using Binary Search: ");
    scanf("%49s", code);
    int index = titans_compare_search_binarysearch(code);
    if (index != -1) {
        printf("Bloom found using Binary Search:\nID:   %d\nCode:   %s\nLevel:
%s\nDescription: %s\n",
            b[index].id, b[index].code, b[index].level, b[index].des);
    } else {
        printf("Bloom with code %s not found using Binary Search.\n", code);
    }
    break;
}
case 6:
    titans_compare_sort_mergesort(0, c - 1);
    printf("Merge Sort complete. Displaying results:\n");
    titans_bloom_retrieve();
    break;
case 7:
    titans_quick_sort(0, c - 1);
    printf("Quick Sort complete. Displaying results:\n");
    titans_bloom_retrieve();
    break;
case 8:
{
    char code[50];
    printf("Enter bloom code to search using Linear Search: ");
    scanf("%49s", code);
    int index = titans_linear_search(code);
    if (index != -1) {
        printf("Bloom found using Linear Search:\nID:   %d\nCode:   %s\nLevel:
%s\nDescription: %s\n",
            b[index].id, b[index].code, b[index].level, b[index].des);
    } else {
        printf("Bloom with code %s not found using Linear Search.\n", code);
    }
    break;
}
case 9:
    titans_complexity_searching();
    break;
case 10:

```

```

        titans_complexity_sorting();
        break;
    case 11:
        printf("Exiting...\n");
        return 0;
    default:
        printf("Invalid choice. Please try again.\n");
        break;
    }
}
}

```

Sorting Algorithms

1. Quick Sort: Algorithm

```

QuickSort(arr[], low, high)
    if low < high
        pivot = Partition(arr[], low, high)
        QuickSort(arr[], low, pivot - 1)
        QuickSort(arr[], pivot + 1, high)

```

```

Partition(arr[], low, high)
    pivot = arr[high]
    i = low - 1
    for j = low to high - 1
        if arr[j] is less than pivot based on the sorting criteria
            swap arr[i] with arr[j]
            i++
    swap arr[i + 1] with arr[high]
    return i + 1

```


Time complexity

Best Case: $O(n \log n)$ (when the pivot divides the array into two equal parts)

Average Case: $O(n \log n)$

Worst Case: $O(n^2)$ (when the pivot is always the smallest or largest element)

- Quick Sort is generally faster on average with $O(n \log n)$ complexity but may degrade to $O(n^2)$ if the pivot selection is poor.

2. Merge Sort:

Algorithm

MergeSort(arr[], left, right)

if left < right

mid = (left + right) / 2

MergeSort(arr[], left, mid)

MergeSort(arr[], mid + 1, right)

Merge(arr[], left, mid, right)

Merge(arr[], left, mid, right)

create left_subarray, right_subarray

copy elements to left_subarray and right_subarray

merge the two subarrays back into arr[] based on the sorting criteria

Time Complexity

Best Case: $O(n \log n)$

Average Case: $O(n \log n)$

Worst Case: $O(n \log n)$ (merge sort always divides the array in half and merges, regardless of the order of elements)

- Merge Sort has a consistent time complexity of $O(n \log n)$, making it reliable for larger datasets or when stability is required.

Searching Algorithms

1. Linear Search:

Algorithm

LinearSearch(arr[], target_code, target_name, target_email)

for each element in arr[]

if element's code == target_code

and element's name == target_name

and element's email == target_email

return index

return -1

Time complexity

- Linear Search is straightforward with $O(n)$ complexity. It doesn't require sorting but is inefficient for larger datasets.

2. Binary Search:

Algorithm

```
BinarySearch(arr[], target_code, target_name, target_email)
    left = 0
    right = arr.length - 1
    while left <= right
        mid = (left + right) / 2
        if arr[mid] matches target based on code, name, and email
            return mid
        else if arr[mid] is less than target based on sorting criteria
            left = mid + 1
        else
            right = mid - 1
    return -1
```

Time Complexity

- Binary Search is efficient with $O(\log n)$ complexity but requires a sorted list. It's well-suited for static datasets where quick lookups are needed.

Comparison of Sorting Algorithms

Quick Sort:

Time Complexity: Average case $\mathcal{O}(n \log n)$; worst case $\mathcal{O}(n^2)$ if the data is already sorted or highly unbalanced when using the last element as a pivot.

Stability: Not a stable sort; may reorder elements with equal keys.

Performance: Fast on average, especially for large, randomized data.

Use Case: Preferred when memory is limited and average case efficiency is prioritized.

Algorithm: Divides data using a pivot element and recursively sorts each partition.

Merge Sort:

Data Requirement: Requires extra memory for temporary arrays but provides stable sorting.

Time Complexity: Consistently $\mathcal{O}(n \log n)$ for all cases, making it reliable.

Stability: Stable sort; maintains order of elements with equal keys.

Performance: Consistent performance across all datasets, ideal for large or pre-sorted data.

Use Case: Ideal when consistent performance and stability are needed, even for sorted or partially sorted data.

Algorithm: Divides data into halves, recursively sorts, and merges them in order.

Comparison of Searching Algorithms

Binary Search:

Data Requirement Requires blooms to be sorted by `bloom code`, `bloom name`, or other relevant fields before searching.

Time Complexity: $O(\log n)$, making it efficient for larger datasets.

Performance: Faster for large, sorted datasets due to logarithmic complexity.

-Use Case: Suitable for quickly finding a specific `bloom code` or other sorted fields in large datasets.

Algorithm Divides the search space in half with each step (binary approach).

Linear Search:

Data Requirement: Works on unsorted data, no pre-sorting needed.

Time Complexity: $O(n)$, as each entry is checked sequentially.

Performance: Slower for large datasets, especially if not sorted.

Use Case: Practical for small or unsorted datasets.

- Algorithm: Checks each bloom entry one-by-one.

SCREEN SHOTS(OUTPUTS)

```

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 1
Enter bloom ID: 1
Enter bloom Code: anxj4431
Enter bloom Level: remembering
Enter bloom Description: retrieve recall or recognise relevant knowledge
Bloom created successfully!

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 1
Enter bloom ID: anxj4561
Invalid input!

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: Invalid input!

```

```

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 1
Enter bloom ID: 2
Enter bloom Code: anxj7433
Enter bloom Level: understand
Enter bloom Description: one or more forms of explanation
Bloom created successfully!

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 1
Enter bloom ID: 3
Enter bloom Code: anxj233
Enter bloom Level: apply
Enter bloom Description: build a new skill
Bloom created successfully!

List of Blooms:
ID: 1
Code: anxj4431
Level: remembering
Description: retrieve recall or recognise relevant knowledge

ID: 2
Code: anxj7433
Level: understand
Description: one or more forms of explanation

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 0
Invalid choice. Please try again.

```

```

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 1
Enter bloom ID: 0
Enter bloom Code: anxj42324
Enter bloom Level: create
Enter bloom Description: put elements together
Bloom created successfully!

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 3

List of Blooms:
ID: 1
Code: anxj4431
Level: remembering
Description: retrieve recall or recognise relevant knowledge

ID: 2
Code: anxj7433
Level: understand
Description: one or more forms of explanation

ID: 3
Code: anxj233
Level: apply
Description: build a new skill

ID: 0
Code: anxj42324
Level: create
Description: put elements together

```

```

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 3

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 7
Quick Sort by ID complete. Displaying results:

```

```

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 5
Enter bloom ID to search using Binary Search: 3
Bloom found using Binary Search:
ID: 3
Code: anxj233
Level: apply
Description: build a new skill

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 2
Enter bloom ID to update: 0
Enter new bloom Code: anxj366
Enter new bloom Level: analyse
Enter new bloom Description: break materials into parts and analyze it
Bloom updated successfully!
1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 9
Linear Search: time complexity is  $O(n)$ .
Binary Search: time complexity is  $O(\log n)$ .

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 10
Quick Sort: time complexity is  $O(n^2)$  for worst case[when last element is chosen as pivot as the elements are already in sorted order] and  $O(n \log n)$  for the average case.
Merge Sort: time complexity is  $O(n \log n)$ .

```

List of Blooms:

ID: 0

Code: anxj42324

Level: create

Description: put elements together

ID: 1

Code: anxj4431

Level: remembering

Description: retrieve recall or recognise relevant knowledge

ID: 2

Code: anxj7433

Level: understand

Description: one or more forms of explanation

ID: 3

Code: anxj233

Level: apply

Description: build a new skill

```
1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 3

List of Blooms:
ID: 0
Code: anxj366
Level: analyse
Description: break materials into parts and analyze it

ID: 1
Code: anxj4431
Level: remembering
Description: retrieve recall or recognise relevant knowledge

ID: 2
Code: anxj7433
Level: understand
Description: one or more forms of explanation

ID: 3
Code: anxj233
Level: apply
Description: build a new skill
```

```
main.c bloom_setting.txt :
1 0 anxj366 analyse break materials into parts and analyze it
2 1 anxj4431 remembering retrieve recall or recognise relevant knowledge
3 2 anxj7433 understand one or more forms of explanation
4 3 anxj233 apply build a new skill
5
```

```
Code: anxj366
Level: analyse
Description: break materials into parts and analyze it

ID: 1
Code: anxj4431
Level: remembering
Description: retrieve recall or recognise relevant knowledge

ID: 2
Code: anxj7433
Level: understand
Description: one or more forms of explanation

ID: 3
Code: anxj233
Level: apply
Description: build a new skill

1. Create Bloom
2. Update Bloom
3. Retrieve Blooms
4. Delete Bloom
5. Search by ID (Binary Search)
6. Sort by ID (Merge Sort)
7. Sort by ID (Quick Sort)
8. Search by ID (Linear Search)
9. Complexity of Searching
10. Complexity of Sorting
11. Exit
Enter your choice: 11
Exiting...
```

Conclusion

The Bloom module, as implemented, provides a comprehensive solution for managing Bloom taxonomy data effectively. The module covers the essential CRUD operations (Create, Retrieve, Update, Delete) while integrating efficient sorting and searching algorithms. By employing both Quick Sort and Merge Sort, the system offers flexibility: Quick Sort is generally faster with average cases, whereas Merge Sort ensures consistent performance and stability.

In addition, the Linear and Binary Search functionalities provide options for both simplicity and efficiency in data retrieval. Linear Search is straightforward and effective for unsorted data, while Binary Search, combined with sorting, enables rapid lookup in sorted datasets. The use of a text file for data storage ensures data persistence and easy retrieval.

Overall, this Bloom module is designed to be robust, efficient, and scalable, making it well-suited for use in educational settings where taxonomy levels and objectives need to be managed systematically. This project demonstrates a balance between simplicity in design and efficiency in execution, meeting the requirements for dynamic data handling and providing a framework adaptable for future enhancements.