

## LAB TASK 11

### **Q1)**

A software team is designing a key-value store using a hash table. The hash table has a size of 7 (indices 0 to 6). The chosen hash function is the Division Method:  $H(key)=key(\text{mod}7)$ .

They need to insert the following sequence of keys: 76, 93, 40, 47, 10, 55.

#### 1. Linear Probing:

- Trace the insertion of all six keys into the hash table using the Linear Probing technique. Assume the probing step is  $(i+1)(\text{mod}7)$ .
- Identify the key(s) that caused the first collision, and state the index where the colliding key was ultimately placed.

#### 2. Separate Chaining:

- If the team had chosen Separate Chaining (Open Hashing) instead, describe the structure of the hash table after all six keys have been inserted.
- For the key 55, state its calculated hash index and describe its final placement in the table under this method.

### **Q2)**

A database system uses a hash table with a current **size of 100** and stores **75 elements**. The hash table uses a collision resolution method that degrades performance significantly when the table becomes too full.

#### 1. Load Factor and Rehashing:

- Calculate the current **Load Factor** for this hash table. Show the formula and the calculated value.
- The system is configured to trigger a **Rehashing** operation when the Load Factor reaches 0.75. Based on your calculation, determine if rehashing is immediately required. Explain your answer.

## 2. Rehashing Implementation:

- If rehashing were to be performed, what is the generally suggested strategy for calculating the **New Table Size**? Propose a new size based on this strategy, starting with the current size of 100.
- Outline the **three main steps** required to execute the rehashing process and replace the old table with the new one

## Q3) String Searching Algorithms

A bioinformatics researcher is analyzing a long DNA sequence (Text, T) to find the locations of a specific gene fragment (Pattern, P).

- Text (T): GCAATGCCCTATGTGACC
- Pattern (P): TATGT

### 1. Brute Force Algorithm:

- Explain the core steps the Brute Force Algorithm takes to find the first occurrence of P in T.
- Trace the first two comparison windows for the pattern starting at Text index 0 and 1, explicitly showing the characters being compared and where the first mismatch occurs for each window.

### 2. Boyer-Moore's Bad Character Heuristic:

- Assume the pattern is aligned starting at T[3]:

T	G	C	A	A	T	G	C	C	T	A	T	G	T	G	A	C	C
P					T	A	T	G	T								

- A mismatch occurs at the last character of the pattern ( $P[4] = 'T'$  vs.  $T[7] = 'C'$ ). The Bad Character is 'C'.
- The pattern P is 'TATGT'. Determine the amount of shift suggested by the Bad Character Heuristic (Case 2: Pattern moves past the mismatch character) in this scenario. Explain the reasoning for the shift distance.

#### **Q4) KMP Prefix Function:**

The Knuth-Morris-Pratt (KMP) Algorithm is known for its linear time complexity in string searching, which is achieved by utilizing a preprocessed structure called the LPS array (Longest Proper Prefix which is also a Suffix).

Consider the following pattern:  $P = \text{"ABABA"}$

##### **1. LPS Array Construction:**

- Construct the LPS array for the pattern  $P = \text{"ABABA"}$ . Show the value for each index.
- State what the LPS value at index 4 (corresponding to the last 'A') means in the context of the KMP algorithm.

##### **2. Skipping Comparisons on Mismatch:**

- Suppose the KMP algorithm is searching for  $\$P\$$  in a text, and a mismatch occurs at the character  $P[4]$  (the last 'A'). The LPS value of the character *before* the mismatch (at  $P[3]$ , the second 'B') is 3.
- Describe the KMP's action following this mismatch. From which index of the pattern will the comparison restart with the current text position?