Comprehensive Guide to Arrays and Strings in Data Structures

Dr. Machbah Uddin

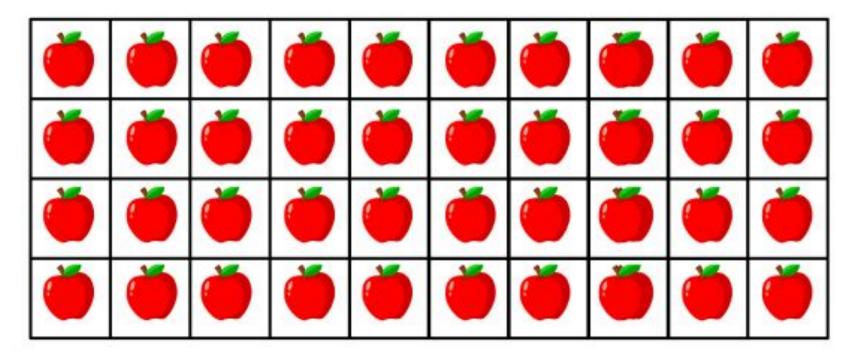
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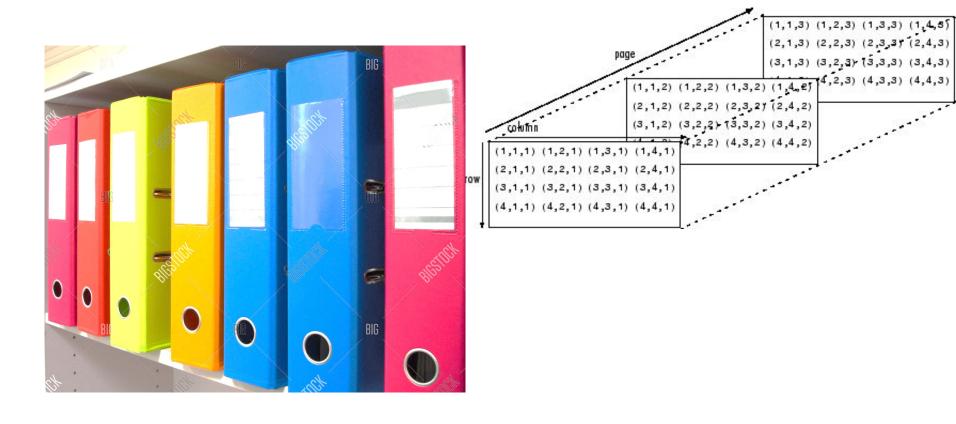
Bangladesh Agricultural University

Arrays

How many apples are there?



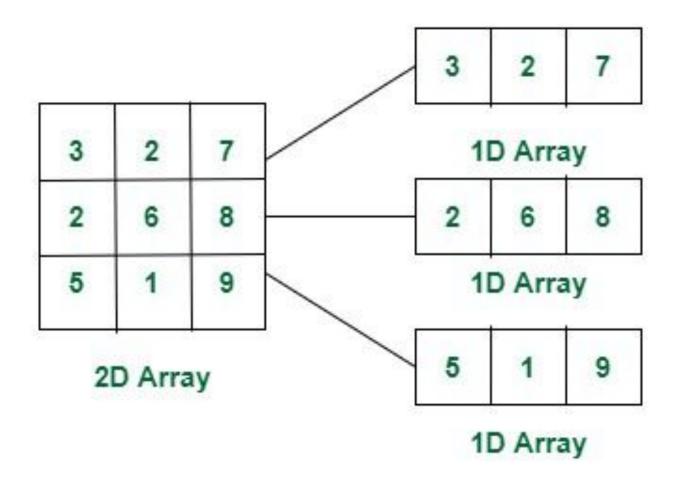
Array



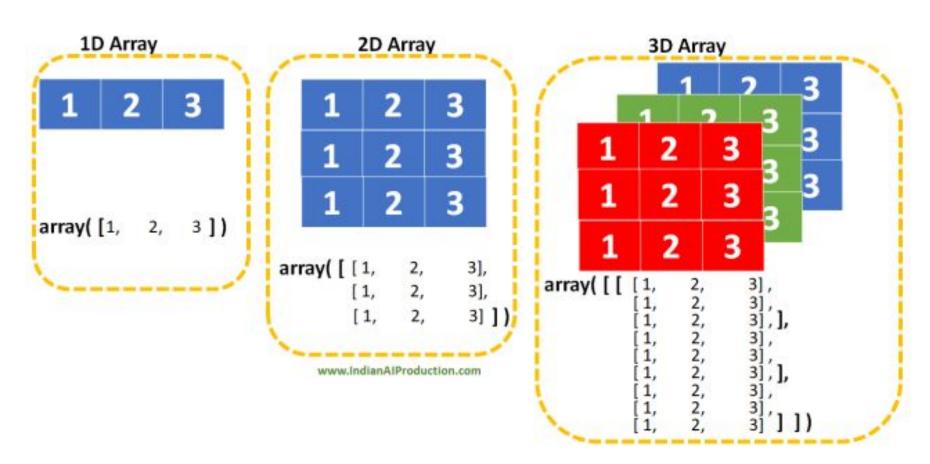
What is an Array?

- An array is a data structure that stores elements of the same type in contiguous memory.
- It allows random access to elements using an index.
- Types: One-dimensional (1D) and Multidimensional Arrays.

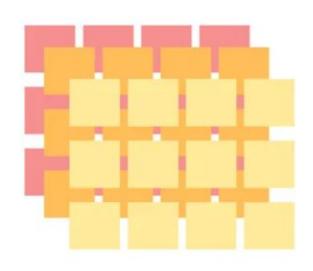
1D and 2D array

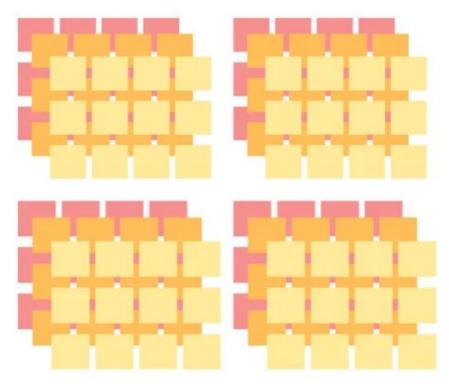


3D arrays



4D arrays





3D array: An array of 2D arrays

4D array: An array of 3D arrays

Why Use Arrays?

- Efficient way to store multiple values.
- Fast access time using indices.
- Saves memory by avoiding overhead of pointers.

One-Dimensional Array (1D)

- A linear data structure where elements are stored in a single row.
- Example: int arr $[5] = \{1, 2, 3, 4, 5\};$

Multidimensional Arrays

- Arrays with more than one index (e.g., 2D, 3D).
- Example (2D Array): int matrix[3][3] = {{1,2,3},{4,5,6},{7,8,9}};

Array Operations Overview

- Insertion
- Traversal
- Deletion
- Searching

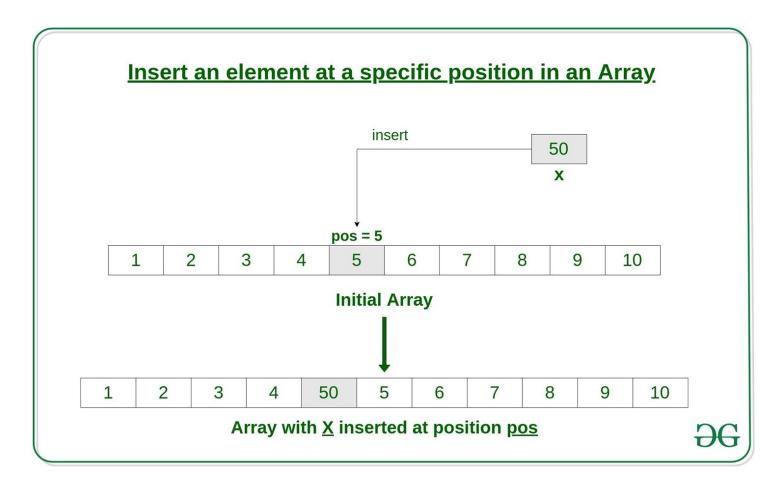
Insertion in Arrays

- Adding an element at a specific position.
- If inserting in the middle, shift elements to the right.

Example: Insertion

- Given array: [10, 20, 30, 40]
- Insert 25 at index 2:
- [10, 20, 25, 30, 40]

Array insert



Traversing an Array

- Accessing each element one by one.
- Can be done using loops.
- Example: for i in range(len(arr)): print(arr[i])

Example: Traversal

• Array: [5, 10, 15, 20]

• Output: 5 10 15 20

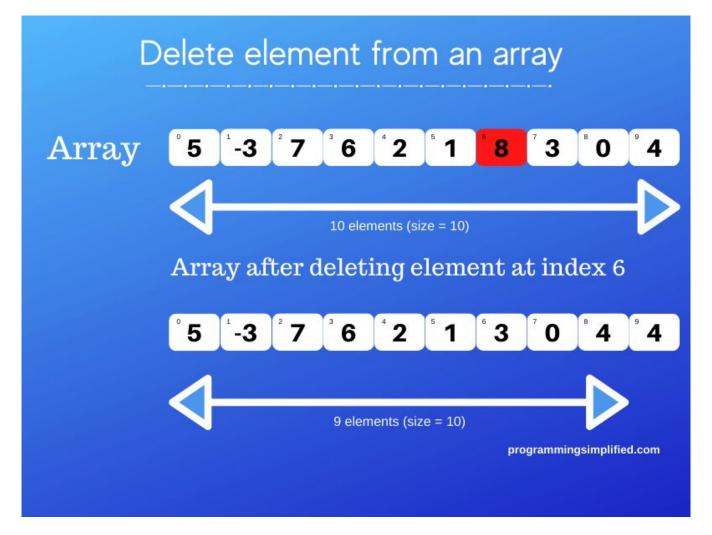
Deletion in Arrays

- Removing an element by shifting elements to the left.
- Last element becomes empty.

Example: Deletion

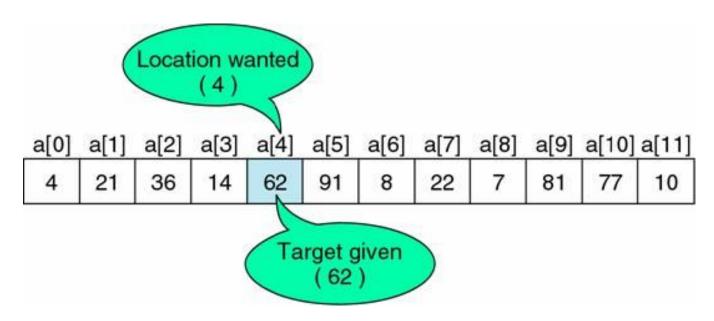
- Given array: [10, 20, 30, 40]
- Delete element at index 2:
- [10, 20, 40]

Array Element Delete



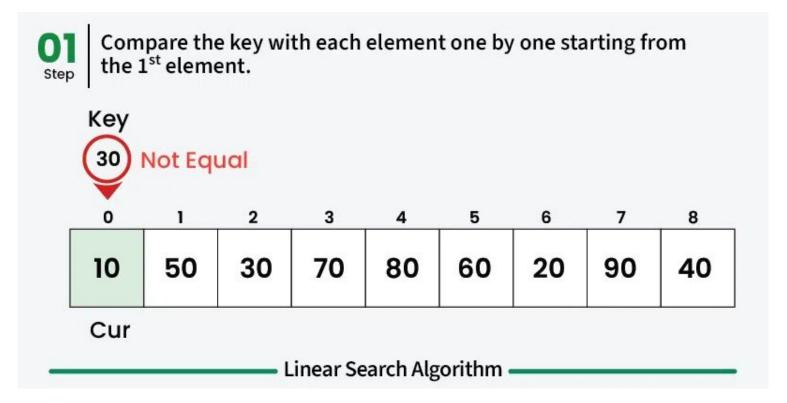
Search Operation in Arrays

- • Linear Search: Check elements one by one.
- Binary Search: Works on sorted arrays, divides array into halves.



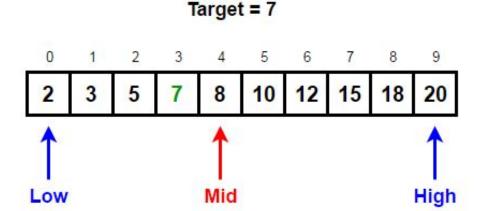
Example: Linear Search

- Array: [5, 10, 15, 20]
- Find $15 \rightarrow$ Found at index 2.



Example: Binary Search

- Sorted Array: [5, 10, 15, 20]
- Find $15 \rightarrow Mid = 10 \rightarrow Search right \rightarrow Found.$



Since 8 (Mid) > 7 (target), we discard the right half and go LEFT

New High = Mid - 1

2D Arrays: Traversal

- Uses nested loops.
- Example:
- for i in range(rows):
- for j in range(cols):
- print(matrix[i][j])

2D Arrays: Insertion

- Specify row and column index.
- Example:
- matrix[1][2] = 9

2D Arrays: Deletion

- Remove an element by shifting elements in a row/column.
- Example:
- Delete row 2 → Shift remaining rows up.

2D Arrays: Searching

- Linear search checks each element.
- Binary search only works on sorted 2D arrays.

Example: 2D Array Search

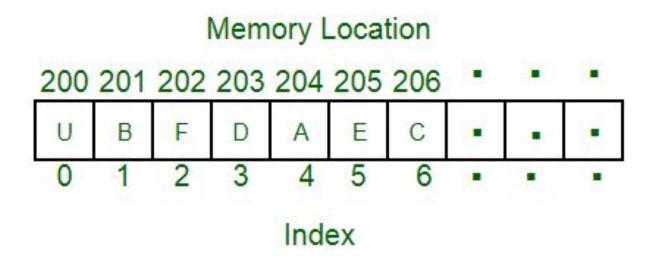
- Given matrix:
- [[1, 2], [3, 4]]
- Find $3 \rightarrow$ Found at (1,0).

Memory Representation of 1D Arrays

- Elements stored in contiguous memory.
- Address calculated as: Base Address + (Index * Size).

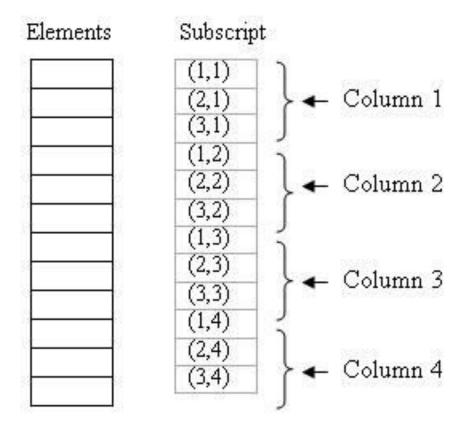
Example: Memory in 1D Arrays

- Array: [10, 20, 30]
- Base Address = 1000, Size = 4 bytes.
- Addresses: 1000, 1004, 1008.



Memory Representation of 2D Arrays

- Stored in Row-Major or Column-Major Order.
- Address = Base + [(Row * ColCount) + Col] *
 ElementSize.



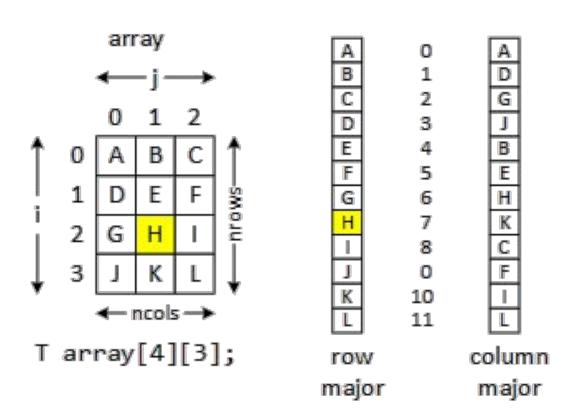
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Example: Memory in 2D Arrays

- Matrix: [[1, 2], [3, 4]]
- Base Address = 1000, Size = 4 bytes.
- Row-Major: 1000, 1004, 1008, 1012.

Comparing Row-Major vs Column-Major

- Row-Major:
 Elements of
 each row stored
 together.
- Column-Major: Elements of each column stored together.

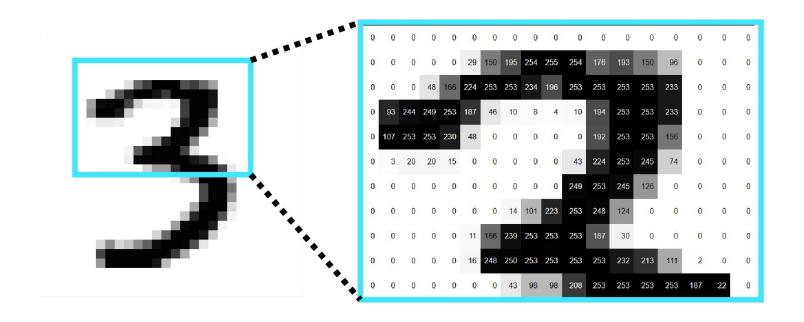


Applications of Arrays

- Used in databases, image processing, and matrices.
- Data storage in programming languages.

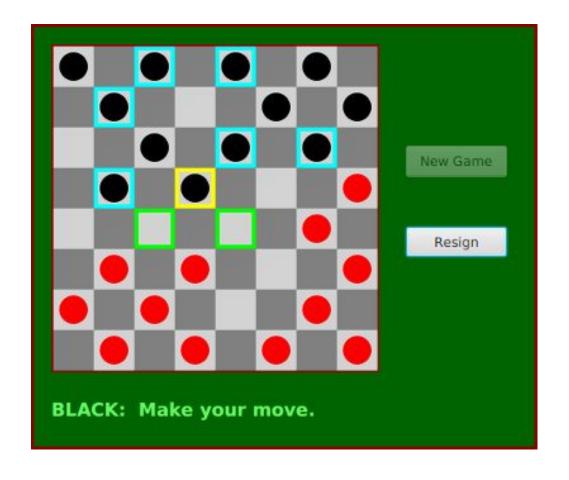
Real-World Example: Image Processing

- Images stored as 2D arrays.
- Pixel values accessed using array indices.



Real-World Example: Game Development

- Grid-based games use 2D arrays.
- Example: Chess board representation.



Challenges with Arrays

- Fixed size (static arrays).
- Costly insertions/deletions in the middle.

Conclusion

- Arrays provide fast access to elements.
- Different operations allow manipulation.
- Used extensively in programming and applications.

Weather Monitoring

- **Problem:** A meteorological department collects hourly temperature readings over a month. Design a system to store and analyze this data (e.g., find the highest, lowest, and average temperatures).
- Why Use an Array?
- The number of readings per day is fixed (24 readings per day \times 30 days = 720 readings).
- Arrays allow fast indexing to access any hour's data efficiently.

Implementation

```
temperatures = [30, 32, 28, 27, 31, 33, 29] #

Example temperatures
highest = max(temperatures)
lowest = min(temperatures)
average = sum(temperatures) / len(temperatures)
print(f"Highest: {highest}, Lowest: {lowest}, Average: {average:.2f}")
```

Stock Price Prediction

- Problem: A stock analyst wants to calculate a
 5-day moving average of stock prices to identify trends.
- Why Use an Array?
- The order of stock prices is crucial (sequential data).
- Arrays provide efficient traversal for calculating moving averages.

Implementation

- stock_prices = [100, 102, 101, 99, 98, 97, 103]
- window_size = 5
- moving_averages =
 [sum(stock_prices[i:i+window_size])/window_
 size for i in
 range(len(stock_prices)-window_size+1)]
- print(moving_averages)

Traffic Management System (Car Plate Number Logs)

- Problem: A smart traffic system stores license plate numbers of 10,000 cars daily to analyze rush hours and detect duplicate entries.
- Why Use an Array?
- The data is **large but structured** (each car plate is stored at a fixed index).
- Arrays allow quick searching and pattern analysis.

Implementation

- car_plates = ["ABC123", "XYZ789", "LMN456", "XYZ789"] # Duplicate detected
- duplicates = set([plate for plate in car_plates if car_plates.count(plate) > 1])
- print(f"Duplicate Plates: {duplicates}")

DNA Sequence Matching

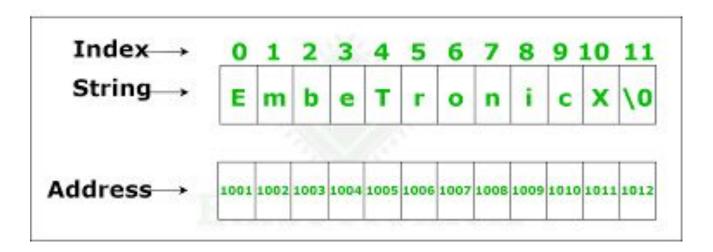
- Problem: Bioinformatics researchers store long DNA sequences as character arrays and need to find whether a smaller DNA sequence exists in a larger one.
- Why Use an Array?
- DNA sequences are naturally stored as arrays of characters.
- Efficient searching algorithms (like KMP or Rabin-Karp) require array-based processing.

Implementation

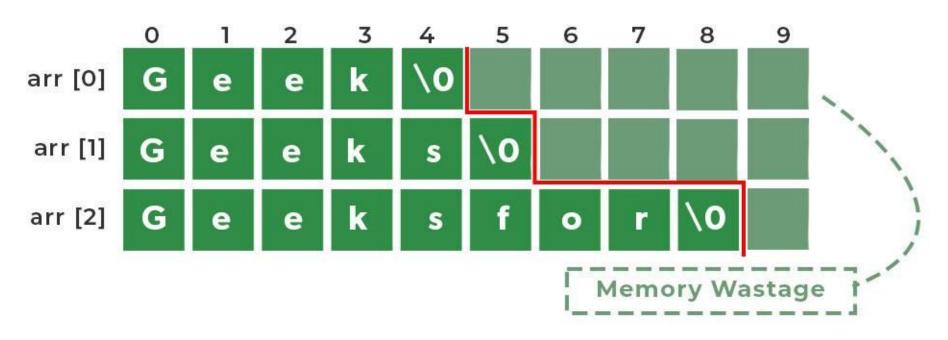
- def is_subsequence(dna, sub):
 return sub in dna
- dna_sequence = "AGCTTAGGCTA"
- sub_sequence = "GGCT"
- print("Found!" if is_subsequence(dna_sequence, sub_sequence) else "Not Found!")

Strings

- Definition: A string is a sequence of characters.
- Examples: "Hello, World!", "DNA Sequence: ATGC", "Binary String: 101010".
- Importance in computer science: Text processing, data storage, and communication.
- Applications: Search engines, bioinformatics, cryptography, and more.



Memory Representation of an Array of Strings



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String Operations

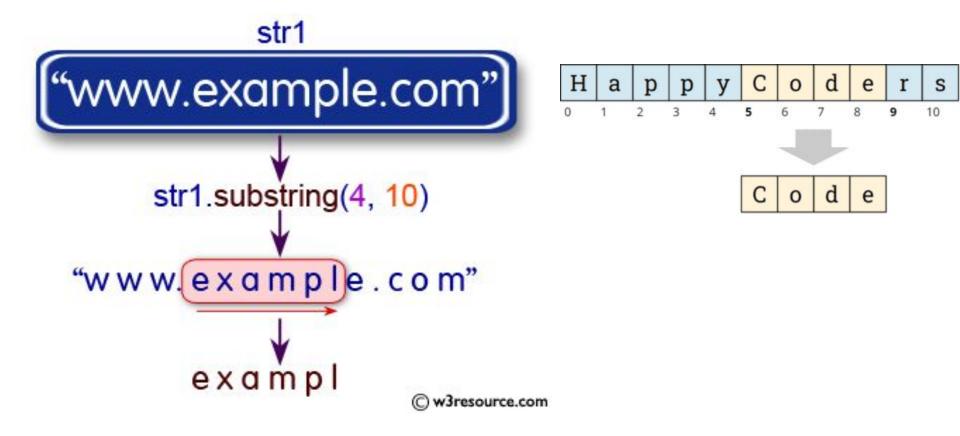
Basic Operations:

- Concatenation: "Hello" + "World" = "HelloWorld".
- Substring extraction:
 "HelloWorld".substring(0, 5) = "Hello".
- Length calculation: len("Hello") = 5.

Advanced Operations:

- Searching for patterns.
- Replacing substrings.
- Splitting and joining strings.

Substring



String searching

Naive (Brute-Force) Search:

- Checks every possible substring.
- Time complexity: O(n*m), where n = text length, m = pattern length.

Knuth-Morris-Pratt (KMP) Algorithm:

- Uses a prefix table to skip unnecessary comparisons.
- Time complexity: O(n + m).

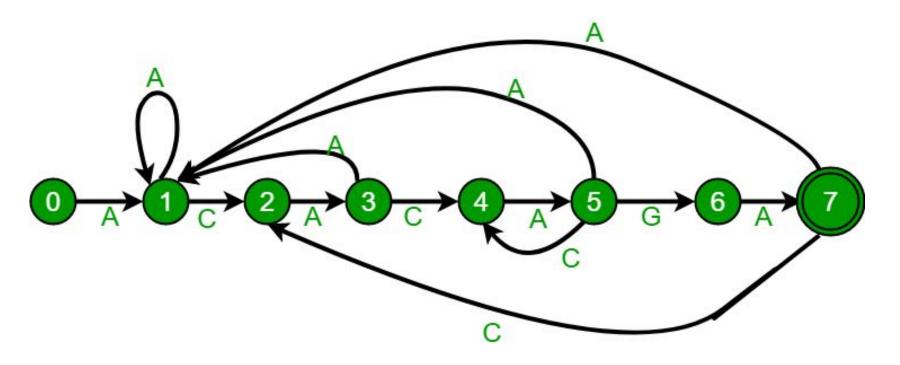
Boyer-Moore Algorithm:

- Skips characters based on bad character and good suffix rules.
- Time complexity: O(n/m) in best case.

Rabin-Karp Algorithm:

- Uses hashing to compare patterns.
- Time complexity: O(n + m) on average.

Pattern Search



Pattern Search

Text: A A B A A C A A D A A B A A B A

Pattern: A A B A

Pattern found at index 0, 9 and 12

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DNA Sequence Matching

- Problem: Given a DNA sequence and a short query sequence, find all occurrences of the query in the DNA sequence.
- Why mandatory? DNA sequences are represented as **strings** (A, T, G, C). Using arrays wouldn't work efficiently for pattern matching.

DNA Sequence Matching

- def find_dna_sequence(dna, query):
 positions = []
 for i in range(len(dna) len(query) + 1):
 if dna[i:i+len(query)] == query:
 positions.append(i)
 return positions
- dna_sequence = "ATGCGATGACCTGAGGCTAGCATGAC"
- query_sequence = "ATG"
- print(find_dna_sequence(dna_sequence, query_sequence)) # Output: [0, 6, 20]

Password Strength Checker

- Problem: Verify if a password meets security standards (length, uppercase, special character, etc.).
- Why mandatory? Passwords are always stored and validated as strings.

Password Strength Checker

- import re
- def check_password_strength(password):
- if (len(password) >= 8 and
- re.search(r'[A-Z]', password) and
- re.search(r'[a-z]', password) and
- re.search(r'[0-9]', password) and
- re.search(r'[@#\$%^&+=]', password)):
- return "Strong Password"
- return "Weak Password"
- print(check_password_strength("Pass@123")) # Output: Strong Password
- print(check_password_strength("pass123")) # Output: Weak Password