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Net Centric Report

Lab: 01

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Problem 1: Hamming

Algorithm

The program would implement two main algorithms:

First, the Hamming Distance function calculates differences between DNA sequences by comparing each position and incrementing a counter whenever characters differ (O(n) time complexity.

- 1. Check if both DNA strings have equal length; if not, panic with error
- 2. Initialize distance counter to zero
- 3. Iterate through each position (i) of both strings simultaneously
- 4. Compare characters at position i in both strings
- 5. If characters differ, increment distance counter
- 6. Return final distance count after completing iteration

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```
func HammingDistance (dna1, dna2 string) int {
    if len(dna1) != len(dna2) {
        panic("DNA sequences must have the same lennth!")
    }

distance := 0
    for i := 0; i < len(dna1); i++ {
        if dna1[i] != dna2[i] {
            distance++
        }
    }

return distance
</pre>
```

Second, the Random DNA generator creates DNA sequences by selecting random nucleotides (A, C, G, T) for each position in a sequence of specified length.

- 1. Define possible nucleotides as "ACGT"
- 2. Seed random number generator with current timestamp
- 3. Create an empty byte slice of requested length
- 4. For each position in the slice:
 - o Generate random number between 0-3
 - Select corresponding nucleotide from "ACGT"
 - Assign selected nucleotide to current position
- 5. Convert completed byte slice to string
- 6. Return generated DNA sequence

```
func RandomDNA(length int) string {

dnaBases := "ACGT"

rand.Seed(time.Now().UnixNano())

dna := make([]byte, length)

for i := 0; i < length; i++ {

dna[i] = dnaBases[rand.Intn(len(dnaBases))]

}

return string(dna)

}
</pre>
```

The program compare two sequences, then generating and comparing 1000 pairs of random DNA sequences.

- 1. Define two example DNA sequences
- 2. Calculate and display the Hamming distance
- 3. Set parameters: 1000 tests with 18-character sequences
- 4. For each test (0 to 999):

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- a. Generate first random DNA sequence
- b. Generate second random DNA sequence
- c. Calculate Hamming distance between sequences
- d. Display test number and distance result
- 5. Display completion message

Output

```
func main() {
    dna1 := "GAGCCTACTAACGGGAT"
    dna2 := "CATCGTAATGACGGCCT"
    fmt.Println("Test with 2 common sample: ")
    fmt.Println("DNA 1:", dna1)
    fmt.Println("DNA 2:", dna2)
    fmt.Println("Hamming Distance: ", HammingDistance(dna1, dna2))

const numTests = 1000
    const dnaLength = 18
    fmt.Println("\nRunning 1000 random DNA tests...")

for i := 0; i < numTests; i++ {
        randDNA1 := RandomDNA(dnaLength)
         result := HammingDistance(randDNA1, randDNA2)
        fmt.Println("Distance ", i, ": ", result)
    }
    fmt.Println("Completed 1000 random DNA tests.")
}</pre>
```

Screenshot 1:

```
• [hopeka@hopeka hamming]$ go run hamming.go
Test with 2 common sample:
DNA 1: GAGCCTACTAACGGGAT
DNA 2: CATCGTAATGACGGCCT
Hamming Distance: 7

Running 1000 random DNA tests...
Distance 0: 14
Distance 1: 11
Distance 2: 12
Distance 3: 14
Distance 4: 18
Distance 5: 15
Distance 6: 15
Distance 7: 11
Distance 8: 14
Distance 9: 11
Distance 9: 11
Distance 9: 11
Distance 10: 13
```

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Screenshot 2:

```
Distance 984 :
               16
Distance 985: 14
Distance 986: 13
Distance 988: 13
Distance 989: 13
Distance 990 : 14
Distance 991: 12
Distance 992: 14
Distance 993: 15
Distance 994: 12
Distance 995: 9
Distance 996: 10
Distance 997: 15
Distance 998: 13
Distance 999: 17
Completed 1000 random DNA tests.
```

Problem 2: Scrabble Score

Algorithm

The program calculates the Scrabble score of a given word.

- 1. Create a map that pass the values to each letter based on Scrabble rules
- 2. Convert the given input string to uppercase to make sure case insensitivity
- 3. Initialize total score, which will store the final calculated score
- 4. Loop through each character in the input string
 - a. If the character exists in the map, retrieve its score

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- b. Add the score to total score
- 5. Return the final score after iterating through all character

Output

```
func main() {

var input string

fmt.Println("Input your word or string for Scrabble Score:")

fmt.Scanln(&input)

score := ScrabbleScore(input)

fmt.Printf("Scrabble Score of'%s' is: %d\n", input, score)

fmt.Printf("Scrabble Score of'%s' is: %d\n", input, score)
```

[hopeka@hopeka scrabble]\$ go run scrabble.go
 Input your word or string for Scrabble Score:
 cabbage
 Scrabble Score of'cabbage' is: 14

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Problem 3: Luhn

Algorithm

```
func isValidLuhn (number string) bool {
   var digits []int
   for _, char := range number {
       if unicode.IsDigit(char) {
           digits = append(digits, int(char-'0'))
   if len(digits) <= 10 {
       fmt.Println("Credit number must be larger than 10")
   sum := 0
    for i := len(digits) - 1; i >= 0; i-- {
       n := digits[i]
       if double {
        sum += n
    return sum%10 == 0
```

The code implements the Luhn algorithm, used for validating credit card numbers.

- 1. Extract all digits from the input string, removing spaces and non-digit characters.
- 2. Check if the number of digits is greater than 10, if not, return false
- 3. Initialize sum to 0 and a boolean flag for altering digits
- 4. Starting from the rightmost digit:
 - a. If the current position needs doubling:
 - i. Double the digit
 - if the doubled value is greater than 0, subtract 9
 - b. Add the resulting digit to the running sum

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- c. Toggle the doubling flag for the next iteration
- 5. Check if the final sum is divisible by 10
- 6. Return true if divisible by 10, false otherwise

Output

```
[hopeka@hopeka Luhn]$ go run luhn.go
4539 3195 0343 6467 - This number is valid!
8273 1232 7352 0569 - This number is not valid!
79927398713 - This number is valid!
```

Problem 4: Minesweeper

Algorithm

The program generates a Minesweeper game board with randomly placed mines and calculates the number of adjacent mines for each empty cell.

- 1. Initialize the Mineflied by creating a 2D board and fillingl all cells with dots to represent empty spaces
- 2. Place the random seed to make sure different placementws each run
- 3. Place the mines in random positions on the board as '*'
 - a. Checking if the cell is not already a mine, place one and increment the mine counter

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```
Codeium: Refactor | Explain | X
func generateMinefield() [] [] rune {

board := make([] [] rune, rows)

for i := range board {

board[i] = make([] rune, cols)

for j := range board[i] {

board[i][j] = '.'

board[i][j] = '.'

}

rand.Seed(time.Now().UnixNano())

for mines := 0; mines < mineCount; {

r, c := rand.Intn(rows), rand.Intn(cols)

if board[r][c] != '*' {

board[r][c] = '*'

mines++

}

return board

return board

return board</pre>
```

- 4. Count adjacent mines for each cell by iterating over the 8 directions and checking if the neighboring cell contains a mine
- Return the total count of mines around the cell

```
func countMines(board [][]rune, r, c int) int {

count := 0

directions := []struct{ dr, dc int }{

{-1, -1}, {-1, 0}, {-1, 1}, {0, -1}, {0, 1}, {1, -1}, {1, 0}, {1, 1},

}

for _, d := range directions {

nr, nc := r+d.dr, c+d.dc

if nr >= 0 && nr < rows && nc >= 0 && nc < cols && board[nr][nc] == '*' {

count++

}

return count

return count
</pre>
```

- 6. Update the board with mine counts by iterating through all cells in the board
 - a. If a cell is empty, call the function countMines to determine the number of adjacent mines
 - b. If at least one mine is found nearby, update the cell with the corresponding number from 1 to 8

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7. Print the final minefield by looping through each row of the board and printing each cell in a formatted way to display the board clearly

```
Codeium: Refactor | Explain | X

func printBoard (board [][]rune) {

for _, row := range board {

for _, cell := range row {

fmt.Printf("%c ", cell)

}

fmt.Println()

}
```

Output

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Problem 5: Matching Brackets

Algorithm

The program checks if a given string contains valid, balanced brackets using a stack.

- 1. Create an empty stack to store opening brackets
- 2. Define a mapping that associates closing brackets with their corresponding
- 3. Loop through each character in the input string
 - a. If the character is an opening bracket push it onto the stack

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- b. if the character is a closing bracket
 - i. Check if the stack is empty, if so, return false
 - ii. Compare the top of stack with the opening bracket using map
 - iii. If they match, pop the top element from the stack, otherwise, return false because of mismatched brackets
- 4. After processing all characters, check if the stack is empty
 - a. If empty, return true as valid
 - b. If not, return false as invalid

Output

```
func main() {

// Các test case

tests := []string{

"([]{})",

"([]]",

""{[()()]}",

""{{[(()()]]}}",

""{{[(()()]]}}",

"""

var tests []string

for _, test := range tests {

fmt.Printf("String \"%s\" -> %v\n", test, isValidBrackets(test))

fmt.Printf("String \"%s\" -> %v\n", test, isValidBrackets(test))

}
```

```
[hopeka@hopeka matching_brackets]$ go run mb.go
String "([]{})" -> true
String "([)]" -> false
String "{[()()]}" -> true
String "{{[(())]]}}" -> true
String "{{[(())]]}}" -> true
String "{[}" -> false
[hopeka@hopeka matching_brackets]$
```