

# OJ review 2

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# Search in Permuted Sorted Array

# Search in Permuted Sorted Array

- Core Idea: BST
  - Split the array into two parts, one part is non-permuted, another one is permuted.
  - Check if the given index is in the non-permuted sorted part
    - If yes, then BST
    - If not, keep splitting

# Find the permutation index

- Initialize:  $l = \text{left}$ ,  $h = \text{right}$
- while( $l < h$ ) {
- $m = (l + h) / 2$
- if( $A[m] > A[h]$ )  $l = m + 1$
- else  $h = m$
- }

Find Repeated DNA Sequences

# Find Repeated DNA Sequences

- Check for every consecutive 10 character-long substring
  - If not in the set
    - Add to the set
  - If exists in the set
    - Output

# Pseudo-code

- Initialize: set{}
- If input in set
  - Output
- Else
  - Set{input}

Find the shortest way



# Find the shortest way

- Open question
  - BFS or DFS
  - Some existing algorithm
    - A\*
    - Dijkstra

Summary of the data structure

# Review of DS

- Assignment 1
  - Overloading
    - Why overload (class operation)
    - How overload (define a member function)
- Assignment 2, 3, 4
  - Binary tree
    - Preorder, postorder, inorder
    - Build heap, priority queue
    - Huffman coding

# Review of DS

- Assignment 5, 6
  - Binary search tree
    - The characteristics of BST (Assignment 5)
    - One typical application (Assignment 6)
- Assignment 7
  - Hash map
- Assignment 8
  - Graph
  - BFS, DFS

# Recommended structure: Binary tree

- Binary tree construction

- 1. Build a node

```
struct node {  
    int data;  
    struct node* left;  
    struct node* right;  
}
```

- 2. Build the tree

```
struct node* NewNode(int data) {  
    struct node* node = new(struct node);  
    node->data = data;  
    node->left = NULL;  
    node->right = NULL;  
  
    return(node);  
}
```

# Recommended structure : Binary tree

- According to your need:
  - Define the member functions
    - `size()`
    - `maxDepth()`
    - `minValue()`
    - `printTree()`
    - ...

# Recommended structure: BST

- BST is built on same node w.r.t. Binary tree

- Difference is the way to insert

```
void BSTTree::Insert(node* & loc,int value)
{
    if (loc == 0)
        struct node* node = new(struct node);
    else if (value<loc->data)
    {
        Insert(loc->left, value);
    }
    else if (value > loc->data)
    {
        Insert(loc->right, value);
    }
    else
        cout << "redundant number!" << endl;
}
```

# Thanks

<http://huangjiezheng.com/teaching/2018-spring-teaching>