

Programming Design

Algorithms and Recursion

Ling-Chieh Kung

Department of Information Management
National Taiwan University

Outline

- **Algorithms and complexity**
- Recursion
- Searching and sorting

Introduction

- It is said that:
 - **Programming** = **Data structures** + **Algorithms**.
 - http://en.wikipedia.org/wiki/Algorithms_%2B_Data_Structures_%3D_Programs
 - To design a program, choose data structures to store your data and choose algorithms to process your data.
- Each of “data structures” and “algorithms” requires one (or more) courses.
 - We will only give you very basic ideas.

Algorithms

- Today we talk about **algorithms**, collections of steps for completing a task.
 - In general, an algorithm is used to **solve a problem**.
 - The most common strategy is to divide a problem into small pieces and then solve those **subproblems**.
 - We will introduce **recursion**, a way to solve a problem based on the solution/outcome of subproblems.
- For a problem, there may be multiple algorithms.
 - The first criterion, of course, is **correctness**.
 - **Time complexity** is typically the next for judging correct algorithms.
- As examples, we introduce two specific problems: **searching** and **sorting**.

Example: listing all prime numbers

- Given an integer n , let's list all the **prime numbers** no greater than n .
- Consider the following (imprecise) algorithm:
 - For each number i no greater than n , check whether it is a prime number.
- To check whether i is a prime number:
 - Idea: If any number $j < i$ can divide i , i is not a prime number.
 - Algorithm: For each number $j < i$, check **whether j divides i** . If there is any j that divides i , report no; otherwise, report yes.
- Before we write a program, we typically prefer to formalize our algorithm.
 - We write **pseudocodes**, a description of steps in words organized in a program structure.
 - This allows us to ignore the details of implementations.

Example: listing all prime numbers

- One pseudocode for listing all prime numbers no greater than n is:

```
Given an integer  $n$ :  
for  $i$  from 2 to  $n$   
    assume that  $i$  is a prime number  
    for  $j$  from 2 to  $i - 1$   
        if  $j$  divides  $i$   
            set  $i$  to be a composite number  
    if  $i$  is still considered as prime  
        print  $i$ 
```

- Implementation:

```
for(int i = 2; i <= n; i++) {  
    bool isPrime = true;  
    for(int j = 2; j < i; j++) {  
        if(i % j == 0) {  
            isPrime = false;  
            break;  
        }  
    }  
    if(isPrime == true)  
        cout << i << " ";  
}
```

- Once we have described an algorithm in pseudocodes, implementation is easy.

A full implementation

- Let's **modularize** our implementation:
 - isPrime(int x)** determines whether the given integer x is a prime number.

```
bool isPrime(int x)
{
    for(int i = 2; i < x; i++)
    {
        if(x % i == 0)
            return false;
    }
    return true;
}
```

- Now we have a correct algorithm.
 - May we improve this algorithm?

```
#include <iostream>
using namespace std;

bool isPrime(int x);
int main()
{
    int n = 0;
    cin >> n;

    for(int i = 2; i <= n; i++)
    {
        if(isPrime(i) == true)
            cout << i << " ";
    }
    return 0;
}
```

Improving our algorithm

- The algorithm can be **faster**:

```
bool isPrime(int x)
{
    for(int i = 2; i * i <= x; i++)
    {
        if(x % i == 0)
            return false;
    }
    return true;
}
```

- Do not use `i <= sqrt(x)` (why?).
 - We improved the algorithm, **not** the implementation.
- May we do even better?

```
#include <iostream>
using namespace std;

bool isPrime(int x);
int main()
{
    int n = 0;
    cin >> n;

    for(int i = 2; i <= n; i++)
    {
        if(isPrime(i) == true)
            cout << i << " ";
    }
    return 0;
}
```


Improving our algorithm further

- Let's consider a completely different algorithm:
 - Let's start from 2. Actually 2, 4, 6, 8, ... are all composite numbers.
 - For 3, actually 3, 6, 9, ... are all composite numbers.
 - We may use a **bottom-up approach** to **eliminate composite numbers**.
- The pseudocode (with comments):

```
Given a Boolean array A of length n
Initialize all elements in A to be true // assuming prime
for i from 2 to n
    if Ai is true
        print i
        for j from 1 to  $\lfloor n/i \rfloor$  // eliminating composite numbers
            Set A[i × j] to false
```

Improving our algorithm further

```
#include <iostream>
using namespace std;

const int MAX_LEN = 10000;

void ruleOutPrime
    (int x, bool isPrime[], int n);

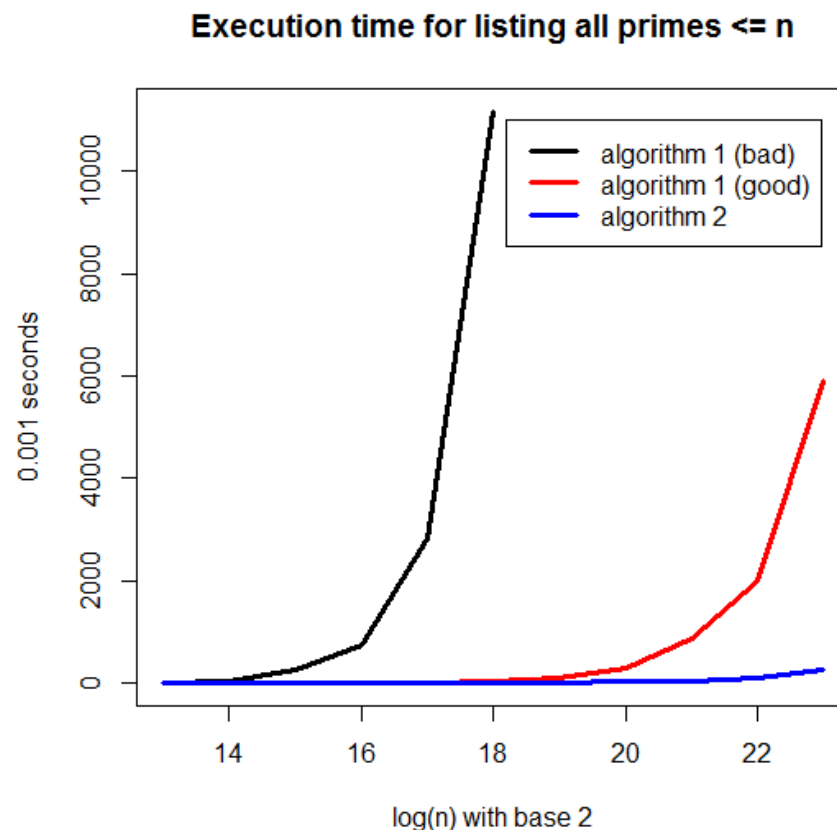
int main()
{
    int n = 0;
    cin >> n; // must < 10000
    bool isPrime[MAX_LEN] = {0};
    for(int i = 0; i < n; i++)
        isPrime[i] = true;
```

```
    for(int i = 2; i <= n; i++)
    {
        if(isPrime[i] == true)
        {
            cout << i << " ";
            ruleOutPrime(i, isPrime, n);
        }
    }
    return 0;
}

void ruleOutPrime
    (int x, bool isPrime[], int n)
{
    for(int i = 1; x * i < n; i++)
        isPrime[x * i] = false;
}
```

Complexity

- While all the three algorithms are correct, they are not equally efficient.
- We typically care about the **complexity** of an algorithm:
 - **Time complexity**: the running time of an algorithm.
 - **Space complexity**: the amount of spaces used by an algorithm.
 - Time is typically more critical.
- Algorithm 2 is much faster!



Complexity

- Running time may be affected by the hardware, number of programs running at the same time, etc.
 - The **number of basic operations** is a better measurement.
 - Basic operations include simple arithmetic, comparisons, etc.
- Convince yourself that algorithm 2 does fewer basic operations.
- The calculation of complexity needs training.
 - This will be formally introduced in Discrete Mathematics, Data Structures, and/or Algorithms.

Outline

- Algorithms and complexity
- **Recursion**
- Searching and sorting

Recursive functions

- A function is **recursive** if it invokes itself (directly or indirectly).
- The process of using recursive functions is called **recursion**.
- Why recursion?
 - Many problems can be solved by dividing the original problem into one or several smaller pieces of **subproblems**.
 - Typically subproblems are **quite similar** to the original problem.
 - With recursion, we write one function to solve the problem by **using the same function** to solve subproblems.

Example 1: finding the maximum

- Suppose that we want to find the maximum number in an array $A[1..n]$ (which means A is of size n).
 - Is there any subproblem whose solution can be utilized?
 - Subproblem: Finding the maximum in an array with size smaller than n .
- A strategy:
 - Subtask 1: First find the maximum of $A[1..(n-1)]$.
 - Subtask 2: Then compare that with $A[n]$.
- How would you visualize this strategy?
- While subtask 2 is simple, subtask 1 is **similar** to the original task.
 - It can be solved with the **same** strategy!

Example 1: finding the maximum

- Let's try to implement the strategy.
- First, I know I need to write a function whose header is:

```
double max(double array[], int len);
```

- This function returns the maximum in **array** (containing **len** elements).
 - I **want** this to happen, though at this moment I do not know how.
- Now let's implement it:
 - If the function **really works**, subtask 1 can be completed by invoking

```
double subMax = max(array, len - 1);
```

- Subtask 2 is done by comparing **subMax** and **array[len - 1]**.

Example 1: finding the maximum

- A (wrong) implementation:
- What will happen if we really invoke this function?
 - The program will not terminate!
 - Even when **len** is 1 in an invocation, we will still try to invoke **max(array, 0)**.
- For an array whose size is 1:
 - That number is the maximum!
- With this, we can add a **stopping condition** into our function.

```
double max(double array[], int len)
{
    double subMax = max(array, len - 1);
    if(array[len - 1] > subMax)
        return array[len - 1];
    else
        return subMax;
}

int main()
{
    double a[5] = {5, 7, 2, 4, 3};
    cout << max(a, 5);
    return 0;
}
```

Example 1: finding the maximum

- A correct implementation is:
- What is the outcome?

```
int main()
{
    double a[5] = {5, 7, 2, 4, 3};
    cout << max(a, 5);
    return 0;
}
```

- Both **else** can be removed. Why?

```
double max(double array[], int len)
{
    if(len == 1) // stopping condition
        return array[0];
    else
    {
        // recursive call
        double subMax = max (array, len - 1);
        if (array[len - 1] > subMax)
            return array[len - 1];
        else
            return subMax;
    }
}
```

Example 1: finding the maximum

- Is it okay to remove both **else**? Why?

```
double max(double array[], int len)
{
    if(len == 1) // stopping condition
        return array[0];
    else
    {
        // recursive call
        double subMax = max (array, len - 1);
        if(array[len - 1] > subMax)
            return array[len - 1];
        else
            return subMax;
    }
}
```

```
double max(double array[], int len)
{
    if(len == 1) // stopping condition
        return array[0];
    // recursive call
    double subMax = max (array, len - 1);
    if(array[len - 1] > subMax)
        return array[len - 1];
    return subMax;
}
```

Example 2: computing factorials

- How to write a function that computes the factorial of n ?
 - A subproblem: computing the factorial of $n - 1$.
 - A strategy: First calculate the factorial of $n - 1$, then multiply it with n .

```
int factorial(int n)
{
    if(n == 1) // stopping condition
        return 1;
    else
        // recursive call
        return factorial(n - 1) * n;
}
```

Example 2: computing factorials

- When we invoke this function with argument 4:
- **factorial(4)**
= **factorial(3) * 4**
= **(factorial(2) * 3) * 4**
= **((factorial(1) * 2) * 3) * 4**
= **((1 * 2) * 3) * 4**
= **(2 * 3) * 4**
= **6 * 4**
= **24**

Example 3: the Fibonacci sequence

- Write a recursive function to find the n th Fibonacci number.
 - The Fibonacci sequence is 1, 1, 2, 3, 5, 8, 13, 21, Each number is the sum of the two proceeding numbers.
 - The n th value can be found once we know the $(n - 1)$ th and $(n - 2)$ th values.

```
int fib(int n)
{
    if(n == 1)
        return 1;
    else if(n == 2)
        return 1;
    else // two recursive calls
        return (fib(n - 1) + fib(n - 2));
}
```

Some remarks

- There must be a **stopping condition** in a recursive function. Otherwise, the program will not terminate.
- In many cases, a recursive strategy can also be implemented with **loops**.
 - E.g., writing a loop for finding a maximum and factorial.
 - But sometimes it is hard to use loops to imitate a recursive function.
- Compared with an equivalent iterative function, a recursive implementation is usually **simpler** and **easier to understand**.
- However, it generally uses **more memory spaces** and is **more time-consuming**.
 - Invoking functions has some cost.

Complexity issue of recursion

- In some cases, recursion is efficient enough.
 - E.g., finding a maximum or calculating the factorial.
- In some cases, however, recursion can be very **inefficient**!
 - E.g., Fibonacci.
- Let's compare the efficiency of two different implementations.

Complexity issue of recursion

- Two implementations:

```
int fib(int n)
{
    if(n == 1)
        return 1;
    else if(n == 2)
        return 1;
    else // two recursive calls
        return (fib(n-1) + fib(n-2));
}
```

```
double fibRepetitive(int n)
{
    if(n == 1 || n == 2)
        return 1;
    int fib1 = 1, fib2 = 1;
    int fib3 = 0;
    for(int i = 2; i < n; i++)
    {
        fib3 = fib1 + fib2;
        fib1 = fib2;
        fib2 = fib3;
    }
    return fib3;
}
```

Complexity issue of recursion

- Which one is faster?

```
int main()
{
    int n = 0;
    cin >> n;
    cout << fibRepetitive(n) << "\n"; // algorithm 1
    cout << fib(n) << "\n"; // algorithm 2
    return 0;
}
```

Polynomial time vs. exponential time

- Given n :
 - The repetitive way has around $c_1 n$ steps, where $c_1 > 0$ is a constant.
 - The recursive way has around $c_2 2^n$ steps, where $c_2 > 0$ is a constant.
- When n is large enough, $c_2 2^n$ is much larger than $c_1 n$.
 - Even if $c_1 \gg c_2$!
 - We say the repetitive way is **more efficient**.
- Technically, we say that:
 - The repetitive way is a **polynomial-time** algorithm
 - The recursive way is an **exponential-time** algorithm.
- In general, an exponential-time algorithm is just too inefficient.

Power of recursion

- Though recursion is sometimes inefficient, typically implementation is easier.
- Let's consider the classic example “**Hanoi Tower**”.
 - There are three pillars and disks of different sizes which can slide onto any pillar. Disc i is smaller than disc j if $i < j$.
 - A large disc cannot be placed on top of a small disc.
- Initially, all discs are at pillar A. We want to move them to pillar C:
 - Only one disk can be moved at a time.
 - Each move consists of taking the upper disk from one of the stacks and placing it on top of another stack.
- What are the steps that solve the Hanoi Tower problem in the fastest way?

A recursive implementation

```
void hanoi(char from, char via,
           char to, int disc)
{
    if(disc == 1)
        cout << "From " << from
              << " to " << to << "\n";
    else
    {
        hanoi(from, to, via, disc - 1);
        cout << "From " << from
              << " to " << to << "\n";
        hanoi(via, from, to, disc - 1);
    }
}
```

```
#include <iostream>
using namespace std;

int main()
{
    int disc = 0; // number of discs
    cin >> disc;
    char a = 'A', b = 'B', c = 'C';

    hanoi(a, b, c, disc);

    return 0;
}
```

- Is there a good way of solving the Hanoi Tower problem iteratively?

Outline

- Algorithms and complexity
- Recursion
- **Searching and sorting**

Searching

- One fundamental task in computation is to **search** for an element.
 - We want to determine whether an element exists in a set.
 - If yes, we want to locate that element.
 - E.g., looking for a string in an article.
- Here we will discuss how to search for an integer in an one-dimensional array.
- Whether the array is **sorted** makes a big difference.

Searching

- Consider an integer array $A[1..n]$ and an integer p .
- How to determine whether p exists in A ?
- If so, where is it?
 - Assume that we only need to find one p even if there are multiple.
- Suppose that the array is unsorted.
- One of the most straightforward way is to apply a **linear search**.
 - Compare each element with p **one by one**, from the first to the last.
 - Whenever we find a match, report its location.
 - Conclude that p does not exist if we end up with nothing.
- The number of operations we need to execute is roughly proportional to n .

Binary search

- What if the array is sorted?
- We may still apply the linear search.
- However, we may improve the efficiency by implementing a **binary search**.
 - First, we compare p with the median m (e.g., $A[(n + 1) / 2]$ if n is odd).
 - If p equals m , bingo!
 - If $p < m$, we know p must exist in **the first half** of A if it exists.
 - If $p > m$, we know p must exist in **the second half** of A if it exists.
 - For the latter two cases, we will continue searching in the **subarray**.

Binary search: pseudocode

```
binarySearch(a sorted array  $A$ , search in between  $from$  and  $to$ , search for  $p$ )  
if  $n = 1$   
    return true if  $A_{from} = p$ ; return false otherwise  
else  
    let  $median$  be floor( $(from + to) / 2$ )  
    if  $p = A_{median}$   
        return true  
    else if  $p < A_{median}$   
        return binarySearch( $A$ ,  $from$ ,  $median$ ,  $p$ )  
    else  
        return binarySearch( $A$ ,  $median + 1$ ,  $to$ ,  $p$ )
```

Linear search vs. binary search

- In binary search, the number of instructions to be executed is roughly proportional to $\log_2 n$.
- So binary search is **much more efficient** than linear search!
 - The difference is huge if the array is large.
 - However, binary search is possible only if the array is sorted.
 - Is it worthwhile to sort an array before we search it?
- It is natural to implement binary search with **recursion**.
 - A subproblem is to search for the element in one half of the array.
- Binary search can also be implemented with repetition.
 - Is it natural to do so?

Sorting

- Given a one-dimensional integer array A of size n , how to sort it?
- Given numbers 6, 9, 3, 4, and 7, how would you sort them?
- Recall what you typically do when you play poker:
 - First put the first number 6 aside.
 - Compare the second number 9 with 6. Because $9 > 6$, put 9 to the right of 6.
 - Compare the third number 3 with the **sorted list** (6, 9). Because $3 < 6$, put 3 to the left of 6.
 - Compare 4 with (3, 6, 9). Because $3 < 4 < 6$, **insert** 4 in between 3 and 6.
 - Compare 7 with (3, 4, 6, 9). Because $6 < 7 < 9$, insert 7 in between 6 and 9.
 - The result is (3, 4, 6, 7, 9).

Insertion sort

- The above algorithm is called **insertion sort**.
 - The key is to maintain a sorted list.
 - Then for each number in the unsorted list, **insert** it into the proper location so that the sorted list **remains sorted**.
- How would you implement the insertion sort?
 - Recursion or repetition?
 - If recursion, what is your strategy?

(Non-repetitive) insertion sort

- The pseudocode:

```
insertionSort(a non-repetitive array  $A$ , the array length  $n$ , an index  $cutoff < n$ )  
// at any time,  $A_{1..cutoff}$  is sorted and  $A_{(cutoff+1)..n}$  is unsorted  
if  $A_{cutoff+1} < A_{1..cutoff}$   
    let  $p$  be 1  
else  
    find  $p$  such that  $A_{p-1} < A_{cutoff+1} < A_p$   
    insert  $A_{cutoff+1}$  to  $A_p$  and shift  $A_{p..cutoff}$  to  $A_{(p+1)..(cutoff+1)}$   
if  $cutoff + 1 < n$   
    insertionSort( $A$ ,  $n$ ,  $cutoff + 1$ )
```

- What if A is repetitive?

Insertion sort

- Roughly how many instructions do we need for insertion sort?
 - We need to do n insertions.
 - To insert the k th value, we search for a position and shift some elements.
 - A linear search: at most k comparisons.
 - Shifting: at most k shifts.
 - Roughly we need $1 + 2 + \cdots + n$ operations, which is proportional to n^2 .
- Does binary search help?

Mergesort (Merge sort)

- Insertion sort is **simple** and fast!
 - Not really “fast”, but faster than many similar sorting algorithm.
 - Because its idea and implementation is simple, it is faster than most algorithms when the array size is **small**.
- Interestingly, there is another sorting algorithm:
 - Its idea is somewhat similar to insertion sort.
 - But it is significantly faster for large arrays!
- This algorithm is called **mergesort**.

Mergesort (Merge sort)

- Recall that in an insertion sort, we need to insert one number into a sorted list for many times.
- A key observation is that “inserting” **another sorted list** of size k into a sorted list can be faster than inserting k separate numbers one by one!
 - So such “inserting” is actually “**merging**”.
- Given an unsorted array, we will:
 - First split the array into two parts, the first half and second half.
 - Then sort each subarray.
 - Finally, merge these two subarrays.
- Mergesort is perfect for recursion!

Mergesort (Merge sort): pseudocode

```
mergeSort(an array  $A$ , the array length  $n$ )  
  let  $median$  be  $\text{floor}((1 + n) / 2)$   
  mergeSort( $A_{1..median}$ ,  $median$ ) // now  $A_{1..median}$  is sorted  
  mergeSort( $A_{(median + 1)..n}$ ,  $n - median + 1$ ) // now  $A_{(median + 1)..n}$  is sorted  
  merge  $A_{1..median}$  and  $A_{(median + 1)..n}$  // how?
```

Mergesort (Merge sort)

- Interestingly, insertion sort is a special way of running mergesort.
 - Not splitting the array into two halves.
 - Instead, splitting it into $A[1..n - 1]$ and $A[n]$.
- Once we use the “smart split”, the **efficiency** is improved a lot!
 - Insertion sort: Roughly proportional to n^2 .
 - Merge sort: Roughly proportional to $n \log n$.
- A simple observation can make a huge difference!