# SELECTION CONTROL OF THE PARTY OF THE PARTY

# Course organization

- Introduction ( Week 1-2)
  - Course introduction
  - A brief introduction to molecular biology
  - A brief introduction to sequence comparison
- Part I: Algorithms for Sequence Analysis (Week 3 8)
  - Chapter 1-3, Models and theories
    - » Probability theory and Statistics (Week 3)
    - » Algorithm complexity analysis (Week 4)
    - » Classic algorithms (Week 5)
  - Chapter 4. Sequence alignment (week 6)
  - Chapter 5. Hidden Markov Models ( week 7 )
  - Chapter 6. Multiple sequence alignment (week 8)
- Part II: Algorithms for Network Biology (Week 9 16)
  - Chapter 7. Omics landscape (week 9)
  - Chapter 8. Microarrays, Clustering and Classification (week 10)
  - Chapter 9. Computational Interpretation of Proteomics (week 11)
  - Chapter 10. Network and Pathways (week 12,13)
  - Chapter 11. Introduction to Bayesian Analysis (week 14,15)
  - Chapter 12. Bayesian networks (week 16)





# Chapter 2: Algorithm Complexity Analysis

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#### **Contents**

- Reading materials
- Why do we need to analyze the complexity of an algorithm?
  - Examples
- Introduction
  - Algorithm complexity
  - "Big O" notation: O()



# Reading

#### Cormen book:

Thomas, H., Cormen, Charles, E., Leiserson, and Ronald, L., Rivest. Introduction to Algorithms, The MIT Press.

(read Chapter 1 and 2, page 1-44).

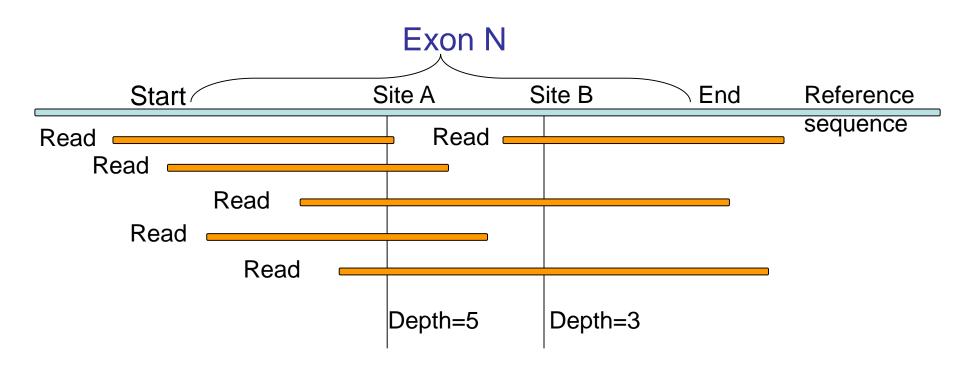


There are ~60 millions of short reads sequenced from exon regions of a human genome. We need to figure out the how many exons were covered with at least 10 reads.

#### Steps:

- 1. Reads are aligned to the genome;
- 2. Each alignment is checked to see the exon it covers;
- 3. For each exon, check the number of reads cover the exon;
- 4. For all exons, filter out those with read number < 10.







#### 1 days later

**Student:** I have created a program to do the analysis. It's running.

Teacher: Cool. Let me know when your analysis finishes.

#### 6 days later...

**Student:** My program has been running for 5 days, and it keeps on running. I have no idea about what is happening and what to do with it.

**Teacher:** Its core is a sorting algorithm with a complexity of at most O(N\*IgN). It should be done within a few minutes!

Student: What?.....



An **algorithm** is any well-defined computational procedure that takes in some **inputs** and produces some **Outputs**.

Example: Sort an array of numbers

 $3, 2, 4, 5, 7, 1, 6 \rightarrow 1, 2, 3, 4, 5, 6, 7$ 



An algorithm is any well-defined computational procedure that takes in some inputs and produces some outputs.

### Complexity: a function of input size

- •Time complexity: the running time
- Space complexity: the memory size required



### Input size

- Number of items in the input
  - Sorting problem
  - FFT
- Total number of bits needed to represent the input
  - $\bullet$ Arithmetic operation (+,-,x,/)
- The value of input
  - Factorial (N!)

### Multiple input sizes

- Need to specify which input size is used
  - Graph operation (number of Vertices, and edges)



#### Before we start

- we use a generic one-processor, randomaccess machine.
  - No parallel



Example: Sort an array of numbers

 $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 



Example: Sort an array of numbers  $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

```
Insertion sort (A)
for j = 2 to length(A)
  do key = A[j]
    /*insert A[j] into the sorted sequence A[1...j-1]
    i=j-1
    while i>0 and A[i]>key
        do A[i+1]=A[i];
        i=i-1;
        A[i+1]=key;
```



```
Example: Sort an array of numbers
            5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6
Insertion sort (A)
for j = 2 to length(A)
     do key = A[j]
          /*insert A[j] into the sorted sequence A[1...j-1]
          i=j-1
          while i>0 and A[i]>key
               do A[i+1]=A[i];
                   i=i-1;
               A[i+1]=key;
```



#### Worst-case and average-case analysis

```
Example: Sort an array of numbers
            5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6
                                           Can repeat
Insertion sort (A)
                                           from 0 to j
for j = 2 to length(A)
                                              times
     do key = A[j]
          /*insert A[j] into the sorted sequence A[1...j-1]
          i=j-1
          while i>0 and A[i]>key
               do A[i+1]=A[i];
                   i=i-1;
               A[i+1]=key;
```



#### Order of growth (增长的阶数)

Example: Sort an array of numbers

 $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

#### Insertion sort:

Algorithm run time complexity: O(N<sup>2</sup>)

Order of growth: 2



# O-notation (big-O notation): Asymptotic upper bound

 $O(g(n)) = \{f(n): \text{ there exist positive constants c}$ and  $n_0$  such that  $0 \le f(n) \le c g(n)$  for all  $n \ge n_0$ 

Note about O-notation operations:  $O(k_1*N^2+k_2*N^3)=O(N^3)$  for constants  $k_1$ ,  $k_2$ 



# O-notation (big-O notation): Asymptotic upper bound

Example: Sort an array of numbers

 $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

Insertion sort: algorithm time complexity: O(N<sup>2</sup>)

#### Sorting with time complexity of O(N\*logN)

Example: Sort an array of numbers  $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

```
Sort (A)
for j = 2 to length(A)

do key = A[j]

/*Use binary search to insert A[j]

/*into the sorted sequence A[1...j-1]

i=j-1
```

Binary\_search(A[j], A[1...j-1],)



Example: Sort an array of numbers

 $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

There are a lot of sorting algorithms:

Heap sort (O(N\*logN))

Merge sort (O(N\*logN))

\*Quick sort (worst-case O(N²), average O(N\*logN))



#### Merge sort

```
Merge-Sort (A, p, r)

if p<r

then q=[(p+r)/2]

Merge-Sort(A, p, q)

Merge-Sort(A,q+1,r)

Merge(A, p, q, r)
```

Time Complexity: 
$$T(N) = \begin{cases} O(1); & \text{if } N = 1\\ 2T(N/2) + O(N); & \text{if } N > 1 \end{cases}$$

Solve it: T(N) = O(N\*logN)



Example: Sort an array of numbers

 $5, 2, 4, 6, 1, 3 \rightarrow 1, 2, 3, 4, 5, 6$ 

Need an array of size N: A[1...N], and 3 temporary variables O(N)

Example: Sequence alignment



#### Examples: Sequence alignment

- Needleman/Wunsch global alignment
- Smith/Waterman local alignment

#### Smith/Waterman local alignment (1981)

- Two sequences  $X = x_1...x_n$  and  $Y = y_1...y_m$
- Let F(i, j) be the optimal alignment score of  $X_{1...i}$  of X up to  $x_i$  and  $Y_{1...j}$  of Y up to  $Y_j$  ( $0 \le i \le n$ ,  $0 \le j \le m$ ), then we have

$$F(0,0) = 0$$

$$F(i, j) = \max \begin{cases} 0 \\ F(i-1, j-1) + S(X_i, Y_j) \\ F(i-1, j) - d \\ F(i, j-1) - d \end{cases}$$



#### Sequence alignment

	Δ	C	A	G	C	C	U	C	G	С	U	U	A	G
Δ	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Α	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0
Α	0.0	0.0	1.0	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.7
U	0.0	0.0	0.0	0.7	0.3	0.0	1.0	0.0	0.0	0.0	1.0	1.0	0.0	0.7
G	0.0	0.0	0.0	1.0	0.3	0.0	0.0	0.7	1.0	0.0	0.0	0.7	0.7	1.0
C	0.0	1.0	0.0	0.0	2.0	1.3	0.3	1.0	0.3	$2 \cdot 0$	0.7	0.3	0.3	0.3
C	0.0	1.0	0.7	0.0	1.0	$3 \cdot 0$	1.7	1.3	1.0	1.3	1.7	0.3	0.0	0.0
Α	0.0	0.0	2.0	0.7	0.3	1.7	2.7	1.3	1.0	0.7	1.0	1.3	1.3	0.0
U	0.0	0.0	0.7	1.7	0.3	1.3	2.7	2.3	1.0	0.7	1.7	2.0	1.0	1.0
U	0.0	0.0	0.3	0.3	1.3	1.0	2.3	2.3	2.0	0.7	1.7	2.7	1.7	1.0
G	0.0	0.0	0.0	1.3	0.0	1.0	1.0	2.0	3.3	2.0	1.7	1.3	$2 \cdot 3$	2.7
Α	0.0	0.0	1.0	0.0	1.0	0.3	0.7	0.7	2.0	3.0	1.7	1.3	$2 \cdot 3$	2.0
C	0.0	1.0	0.0	0.7	1.0	2.0	0.7	1.7	1.7	3.0	2.7	1.3	1.0	$2 \cdot 0$
G	0.0	0.0	0.7	1.0	0.3	0.7	1.7	0.3	2.7	1.7	2.7	$2 \cdot 3$	1.0	$2 \cdot 0$
G	0.0	0.0	0.0	1.7	0.7	0.3	0.3	1.3	1.3	$2 \cdot 3$	1.3	$2 \cdot 3$	$2 \cdot 0$	$2 \cdot 0$

Fig. 1.  $H_{ij}$  matrix generated from the application of eqn (1) to the sequences A-A-U-G-C-C-A-U-U-G-A-C-G-G and C-A-G-C-C-U-C-G-C-U-U-A-G. The underlined elements indicate the trackback path from the maximal element 3·30.

Smith and Waterman, JMB, 1981, 147, 195-197

Time complexity: O(N\*M)

Space complexity: O(N\*M) or O(max(N, M))

Need a two-dimension array of size N\*M, and a constant number of temporary variables



### Other issues impact the speed of a program

- Output size
  - Blast: output can be a problem
- Input/Output method/place/mode
  - Speed
    - screen << hard disk << memory</li>
- Programming language
  - Speed
    - ●Perl < java < C++ <C