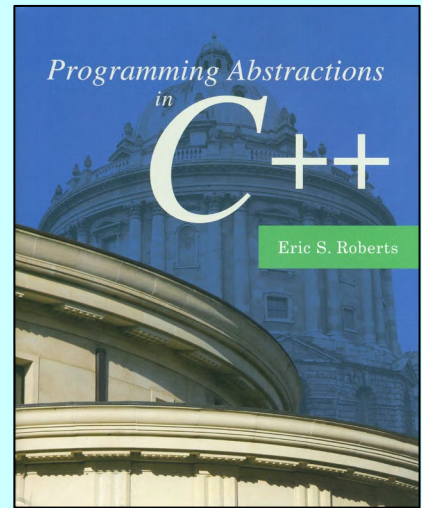


## CHAPTER 10

# Algorithmic Analysis

*Without analysis, no synthesis.*

—Friedrich Engels, *Herr Eugen Dühring's  
Revolution in Science*, 1878



10.1 The sorting problem: selection sort

10.2 Computational complexity

10.3 Recursion to the rescue: merge sort

10.4 Standard complexity classes

10.5 The Quicksort algorithm

# The Sorting Problem

- Of all the algorithmic problems that computer scientists have studied, the one with the broadest practical impact is certainly the *sorting* problem, which is the problem of arranging the elements of an array or a vector in order.

vec							
56	25	37	58	95	19	73	30
0	1	2	3	4	5	6	7
19	25	30	37	56	58	73	95
0	1	2	3	4	5	6	7

- The sorting problem comes up, for example, in alphabetizing a telephone directory, arranging library records by catalogue number, and organizing a bulk mailing by ZIP code.
- There are many algorithms that one can use to sort an array. Because these algorithms vary enormously in their efficiency, it is critical to choose a good algorithm, particularly if the application needs to work with large arrays.



# Visualizing Sorting Algorithms

[www.cs.usfca.edu/~galles/visualization/ComparisonSort.html](http://www.cs.usfca.edu/~galles/visualization/ComparisonSort.html)

[www.toptal.com/developers/sorting-algorithms](http://www.toptal.com/developers/sorting-algorithms)

 Play All	 Insertion	 Selection	 Bubble	 Shell	 Merge	 Heap	 Quick	 Quick3
 Random								
 Nearly Sorted								
 Reversed								
 Few Unique								

# The Selection Sort Algorithm

- Of the many sorting algorithms, the easiest one to describe is *selection sort*, which appears in the text like this:

```
void sort(Vector<int> & vec) {  
    int n = vec.size();  
    for (int lh = 0; lh < n; lh++) {  
        int rh = lh;  
        for (int i = lh + 1; i < n; i++) {  
            if (vec[i] < vec[rh]) rh = i;  
        }  
        int temp = vec[lh];  
        vec[lh] = vec[rh];  
        vec[rh] = temp;  
    }  
}
```

- Coding this algorithm as a single function makes sense for efficiency but complicates the analysis. The next two slides decompose selection sort into a set of functions that make the operation easier to follow.

# Decomposition of the `sort` Function

```
/*
 * Function: sort
 * -----
 * Sorts a Vector<int> into increasing order. This implementation
 * uses an algorithm called selection sort, which can be described
 * in English as follows. With your left hand (lh), point at each
 * element in the vector in turn, starting at index 0. At each
 * step in the cycle:
 *
 * 1. Find the smallest element in the range between your left
 *    hand and the end of the vector, and point at that element
 *    with your right hand (rh).
 *
 * 2. Move that element into its correct position by swapping
 *    the elements indicated by your left and right hands.
 */

void sort(Vector<int> & vec) {
    for ( int lh = 0 ; lh < vec.size() ; lh++ ) {
        int rh = findSmallest(vec, lh, vec.size() - 1);
        swap(vec[lh], vec[rh]);
    }
}
```

# Decomposition of the `sort` Function

```
/*  
 * Function: findSmallest  
 * -----  
 * Returns the index of the smallest value in the vector between  
 * index positions p1 and p2, inclusive.  
 */
```

```
int findSmallest(Vector<int> & vec, int p1, int p2) {  
    int smallestIndex = p1;  
    for ( int i = p1 + 1 ; i <= p2 ; i++ ) {  
        if (vec[i] < vec[smallestIndex]) smallestIndex = i;  
    }  
    return smallestIndex;  
}
```

```
/*  
 * Function: swap  
 * -----  
 * Exchanges two integer values passed by reference.  
 */
```

```
void swap(int & x, int & y) {  
    int temp = x;  
    x = y;  
    y = temp;  
}
```

# Simulating Selection Sort

```
int main() {
```

```
    void sort(Vector<int> & vec) {
```

```
        for ( int lh = 0 ; lh < vec.size() ; lh++ ) {
```

```
            int rh = findSmallest(vec, lh, vec.size() - 1);
```

```
            swap(vec[lh], vec[rh]);
```

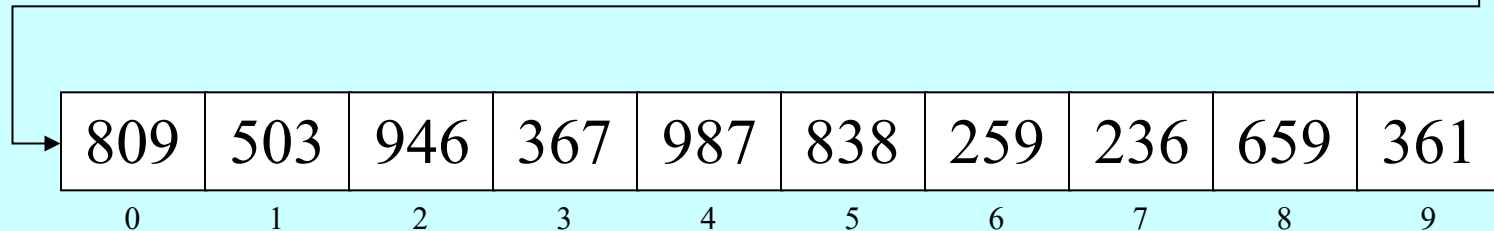
```
        }
```

```
    }
```

lh

rh

vec



# Efficiency of Selection Sort

- The primary question for today is how one might evaluate the efficiency of an algorithm such as selection sort.
- One strategy is to measure the **actual** time it takes to run for arrays of different sizes. In C++, you can measure elapsed time by calling the **time** function, which returns the current time in milliseconds.

```
#include <time.h>
/* time_t, struct tm, difftime, time, mktime */
int main() {
    Vector<int> vec = createTestVector();
    time_t begin, end;
    time(&begin);
    sort(vec);
    double diff = difftime(time(&end), begin);
    return 0;
}
```



# Efficiency of Selection Sort

- Using this strategy (measuring the **actual** running time), however, requires some care:
  - The **time** function is often too rough for accurate measurement. It therefore makes sense to measure **several runs** together and then divide the total time by the number of repetitions.
  - Most algorithms show some variability depending on the data. To avoid distortion, you should run **several independent trials** with different data and average the results.
  - Some measurements are likely to be **wildly off** because the computer needs to run some background task. Such data points must **be discarded** as you work through the analysis.

# Measuring Sort Timings

The following table shows the average timing of the selection sort algorithm after removing outlying trials that differ by more than two standard deviations from the mean. The column labeled  $\mu$  (the Greek letter *mu*, which is the standard statistical symbol for the mean) is a reasonably good estimate of running time.

	Trial 1	Trial 2	Trial 3	Trial 4	Trial 5	Trial 6	Trial 7	Trial 8	Trial 9	Trial 10	$\mu$	$\sigma$
$N = 10$	.0021	.0025	.0022	.0026	.0020	.0030	.0022	.0023	.0022	.0025	.0024	.00029
20	.006	.007	.008	.007	.007		.007	.007	.007	.007	.007	.00036
30	.014	.014	.014	.015	.014	.014	.014	.014	.014	.014	.014	.00013
40	.028	.024	.025	.026	.023	.025	.025	.026	.025	.027	.025	.0014
50	.039	.037	.036	.041	.042	.039		.039	.034	.038	.039	.0025
100	.187	.152	.168	.176	.146	.146	.165	.146	.178	.154	.162	.0151
500	3.94	3.63	4.06	3.76	4.11	3.51	3.48	3.64	3.31	3.45	3.69	0.272
1000	13.40	12.90	13.80	17.60	12.90	14.10	12.70		16.00	15.50	14.32	1.69
5000	322.5	355.9	391.7	321.6	388.3	321.3	321.3	398.7	322.1	321.3	346.4	33.83
10000	1319.		1327.	1318.	1331.	1336.	1318.	1335.	1325.	1319.	1326.	7.50

# Selection Sort Running Times

- Many algorithms that operate on vectors have running times that are proportional to the size of the array. If you multiply the number of values by ten, you would expect those algorithms to take ten times as long.
- As the running times on the preceding slide make clear, the situation for selection sort is very different. The table on the right shows the average running time when selection sort is applied to 10, 100, 1000, and 10000 values.

<i>N</i>	<i>time</i>
10	.0024
100	0.162
1000	14.32
10000	1332.

- As a rough approximation—particularly as you work with larger values of  $N$ —it appears that every ten-fold increase in the size of the array means that selection sort takes about 100 times as long.

# Counting Operations

- Another way to estimate the running time is to count how many operations are required to sort an array of size  $N$ .
- In the selection sort implementation, the section of code that is executed most frequently (and therefore contributes the most to the running time) is the body of the **findSmallest** method. The number of operations involved in each call to **findSmallest** changes as the algorithm proceeds:
  - $N$  values are considered on the first call to **findSmallest**.
  - $N - 1$  values are considered on the second call.
  - $N - 2$  values are considered on the third call, and so on.
- In mathematical notation, the number of values considered in **findSmallest** can be expressed as a summation, which can then be transformed into a simple formula:

$$1 + 2 + 3 + \cdots + (N - 1) + N = \sum_{i=1}^N i = \frac{N \times (N + 1)}{2}$$

# Quadratic Growth

- The reason behind the rapid growth in the running time of selection sort becomes clear if you make a table showing the value of  $N(N+1)/2$  for various values of  $N$ :

$N$	$\frac{N \times (N + 1)}{2}$
10	55
100	5050
1000	500,500
10000	50,005,000

- The growth pattern in the right column is similar to that of the measured running time of the selection sort algorithm. As the value of  $N$  increases by a factor of 10, the value of  $N(N+1)/2$  increases by a factor of around 100, which is  $10^2$ . Algorithms whose running times increase in proportion to the square of the problem size are said to be *quadratic*.



# Big-O Notation

- The most common way to express computational complexity is to use ***big-O notation***, which was introduced by the German mathematician Paul Bachmann in 1892.
- Big-O notation consists of the letter  $O$  followed by a formula that offers a qualitative assessment of running time as a function of the problem size, traditionally denoted as  $N$ . E.g., the computational complexity of linear search is  $O(N)$  and the computational complexity of selection sort is  $O(N^2)$ .
- More formally, if  $f(N) = O(g(N))$ ,  $\exists C > 0, \exists N_0, \forall N > N_0$ ,  $f(N) \leq C g(N)$ .
- If you read these formulas aloud, you would pronounce them as “big-O of  $N$ ” and “big-O of  $N^2$ ” respectively.



# Common Simplifications of Big-O

- Given that big-O notation is designed to provide a qualitative assessment, it is important to make the formula inside the parentheses as simple as possible.
- When you write a big-O expression, you should always make the following simplifications:
  1. Eliminate any term whose contribution to the running time ceases to be significant as  $N$  becomes large.
  2. Eliminate any constant factors.
- The computational complexity of selection sort is therefore

$$O(N^2)$$

and not

$$O\left(\frac{N \times (N+1)}{2}\right)$$



# Deducing Complexity from the Code

- In many cases, you can deduce the computational complexity of a program directly from the structure of the code.
- The standard approach to doing this type of analysis begins with looking for any section of code that is executed more often than other parts of the program. As long as the individual operations involved in an algorithm take roughly the same amount of time, the operations that are executed most often will come to dominate the overall running time.
- In the selection sort implementation, for example, the most commonly executed statement is the `if` statement inside the `findSmallest` method. This statement is part of two `for` loops, one in `findSmallest` itself and one in `Sort`. The total number of executions is

$$1 + 2 + 3 + \cdots + (N - 1) + N$$

which is  $O(N^2)$ .





# Worst-case vs. average-case complexity

- When you analyze the computational complexity of a program, you're usually not interested in the minimum possible time (*Best-case complexity*) but, in general, the following two types of complexity analysis:
  - *Worst-case complexity*: The upper bound on the computational complexity. You can *guarantee* that the performance of the algorithm will be at least as good as your analysis indicates.
  - *Average-case complexity*: The best statistical estimate of actual performance. Usually much more difficult to carry out and typically requires considerable mathematical sophistication.
- E.g., for linear search:
  - the worse-case complexity is  $O(N)$
  - the average-case complexity is  $O(N)$
  - the best-case complexity is  $O(1)$

# Worst-case is more interesting

- The worst-case running time of an algorithm gives us an upper bound on the running time for *any input*. Knowing it provides a guarantee that the algorithm will never take any longer. We need not make some educated guess about the running time and hope that it never gets much worse.
- For some algorithms, the worst case occurs fairly often. E.g., in searching a database for a particular piece of information, the searching algorithm's worst case will often occur when the information is not present in the database. In some applications, searches for absent information may be frequent.
- The average case is often roughly as bad as the worst case. E.g., in linear search, on average, we check half of the elements. The resulting average-case running time turns out to be a linear function of the input size, just like the worst-case running time.

-- *Introduction to Algorithms*, CLRS.

# Recursion to the Rescue

- As long as arrays are small, selection sort is a perfectly workable strategy. Even for 10,000 elements, the average running time of selection sort is just over a second.
- The quadratic behavior of selection sort, however, makes it less attractive for the very large arrays that one encounters in commercial applications. Assuming that the quadratic growth pattern continues beyond the timings reported in the table, sorting 100,000 values would require two minutes, and sorting 1,000,000 values would take more than three hours.
- The computational complexity of the selection sort algorithm, however, holds out some hope:
  - Sorting twice as many elements takes four times as long.
  - **Sorting half as many elements takes only one fourth the time.**
  - Is there any way to use sorting half an array as a subtask in a recursive solution to the sorting problem?

# The Merge Sort Idea

1. Divide the vector into two halves: **v1** and **v2**.
2. Sort each of **v1** and **v2** recursively.
3. Clear the original vector.
- 4 Merge elements into the original vector by choosing the smallest element from **v1** or **v2** on each cycle.

**vec**

236	259	361	367	503	659	809	838	946	987
0	1	2	3	4	5	6	7	8	9

**v1**

0	1	2	3	4

**v2**

0	1	2	3	4

# The Merge Sort Implementation

```
/*  
 * The merge sort algorithm consists of the following steps:  
 *  
 * 1. Divide the vector into two halves.  
 * 2. Sort each of these smaller vectors recursively.  
 * 3. Merge the two vectors back into the original one.  
 */  
  
void sort(Vector<int> & vec) {  
    int n = vec.size();  
    if (n <= 1) return;  
    Vector<int> v1;  
    Vector<int> v2;  
    for (int i = 0; i < n; i++) {  
        if (i < n / 2) {  
            v1.add(vec[i]);  
        } else {  
            v2.add(vec[i]);  
        }  
    }  
    sort(v1);  
    sort(v2);  
    vec.clear();  
    merge(vec, v1, v2);  
}
```

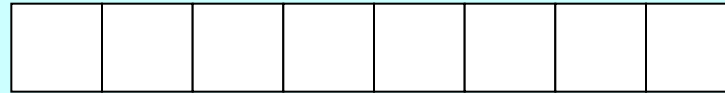
# The Merge Sort Implementation

```
/*
 * Function: merge
 * -----
 * This function merges two sorted vectors (v1 and v2) into the
 * vector vec, which should be empty before this operation.
 * Because the input vectors are sorted, the implementation can
 * always select the first unused element in one of the input
 * vectors to fill the next position.
 */

void merge(Vector<int> & vec, Vector<int> & v1, Vector<int> & v2) {
    int n1 = v1.size();
    int n2 = v2.size();
    int p1 = 0;
    int p2 = 0;
    while (p1 < n1 && p2 < n2) {
        if (v1[p1] < v2[p2]) {
            vec.add(v1[p1++]);
        } else {
            vec.add(v2[p2++]);
        }
    }
    while (p1 < n1) vec.add(v1[p1++]);
    while (p2 < n2) vec.add(v2[p2++]);
}
```

# The Complexity of Merge Sort

Sorting 8 items



*requires*

Two sorts of 4 items



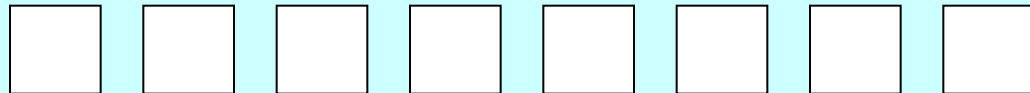
*which requires*

Four sorts of 2 items



*which requires*

Eight sorts of 1 item



The work done at each level (*i.e.*, the sum of the work done by all the calls at that level) is proportional to the size of the vector.

The running time is therefore proportional to  $N$  times the number of levels.

# How Many Levels Are There?

- The number of levels in the merge sort decomposition is equal to the number of times you can divide the original vector in half until there is only one element remaining. In other words, what you need to find is the value of  $k$  that satisfies the following equation:

$$1 = N / \underbrace{2 / 2 / 2 / 2 \cdots / 2}_{k \text{ times}}$$

- You can simplify this formula using basic mathematics:

$$1 = N / 2^k$$

$$2^k = N$$

$$k = \log_2 N$$

- The complexity of merge sort is therefore  $O(M \log N)$ .



# Comparing $N^2$ and $N \log N$

- The difference between  $O(N^2)$  and  $O(N \log N)$  is enormous for large values of  $N$ , as shown in this table:

$N$	$N^2$	$N \log_2 N$
10	100	33
100	10,000	664
1,000	1,000,000	9,966
10,000	100,000,000	132,877
100,000	10,000,000,000	1,660,964
1,000,000	1,000,000,000,000	19,931,569

- Based on these numbers, the theoretical advantage of using merge sort over selection sort on a vector of 1,000,000 values would be a factor of more than 50,000.



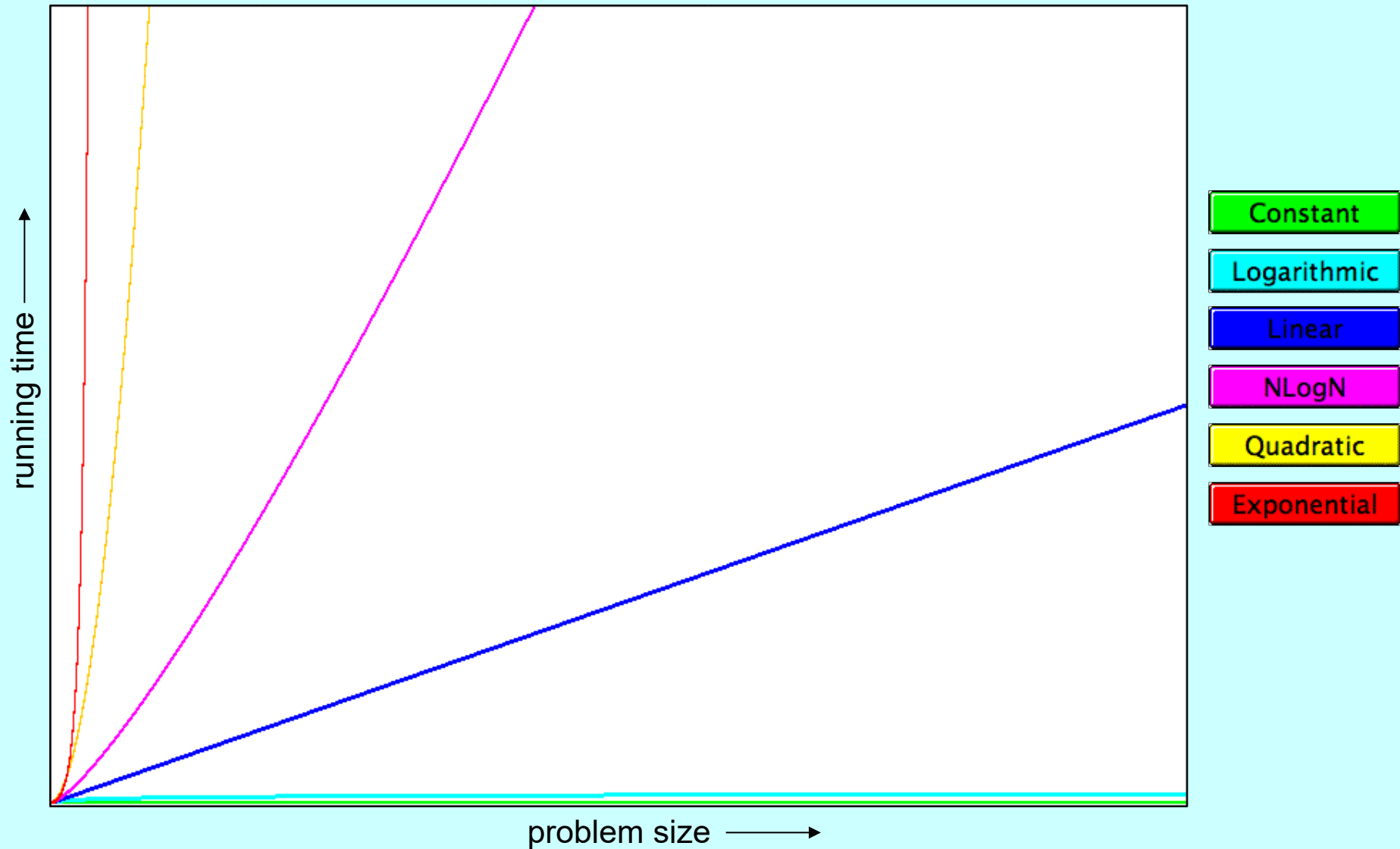
# Standard Complexity Classes

- The complexity of a particular algorithm tends to fall into one of a small number of standard complexity classes:

Constant	$O(1)$	$O(\textit{Yeah})$ Finding elements in a vector using indices
Logarithmic	$O(\log N)$	$O(\textit{Nice})$ Binary search in a sorted vector
Linear	$O(N)$	$O(\textit{Fine})$ Summing a vector; linear search
$M\log N$	$O(M\log N)$	$O(K)$ Merge sort
Quadratic	$O(N^2)$	$O(\textit{Well})$ Selection sort
Cubic	$O(N^3)$	$O(\textit{My})$ Obvious algorithms for matrix multiplication
Exponential	$O(2^N)$	$O(\textit{No})$ Tower of Hanoi solution
Factorial	$O(N!)$	$O(\textit{MG})$ Brute-force search through all permutations

- In general, theoretical computer scientists regard any problem whose complexity cannot be expressed as a polynomial as *intractable*.










# Graphs of the Complexity Classes



# The Quicksort Algorithm

- Most sorting libraries use some variation of the *Quicksort algorithm*, which was developed by Tony Hoare in the 1950s.
- The Quicksort algorithm consists of two phases:
  1. **Partition.** In the partition phase, the elements of the array are reordered so that the array begins with a set of “small” elements and ends with a set of “big” elements, where the distinction between “small” and “big” is made relative to an element called the *pivot*, which appears at the boundary between the two regions.
  2. **Sort.** In the sort phase, the Quicksort algorithm is applied recursively to the “small” and “big” subarrays, which leaves the entire array sorted.

# Partitioning the Array

<del>809</del>	503	<del>946</del>	367	<del>089</del>	<del>838</del>	<del>809</del>	<del>838</del>	<del>089</del>	<del>946</del>
0	1	2	3	4	5	6	7	8	9
									

1. Select the first element as the ***pivot*** and set it aside.
2. Keep two indices into the remaining elements, starting at each end.
3. Advance the indices until the left index is larger than the pivot and the right index is smaller.
4. Exchange the elements at the two indices.
5. Repeat the process until the indices coincide.
6. Swap the pivot and the index.

# The Quicksort Implementation

```
/*
 * Function: sort
 * -----
 * In this implementation, sort is a wrapper function that
 * calls quicksort to do all the work.
 */
void sort(Vector<int> & vec) {
    quicksort(vec, 0, vec.size() - 1);
}

/*
 * Function: quicksort
 * -----
 * Sorts the elements in the vector between index positions
 * start and finish, inclusive. The Quicksort algorithm begins
 * by "partitioning" the vector so that all elements smaller
 * than a designated pivot element appear to the left of a
 * boundary and all equal or larger values appear to the right.
 * Sorting the subsidiary vectors to the left and right of the
 * boundary ensures that the entire vector is sorted.
 */
void quicksort(Vector<int> & vec, int start, int finish) {
    if (start >= finish) return;
    int boundary = partition(vec, start, finish);
    quicksort(vec, start, boundary - 1);
    quicksort(vec, boundary + 1, finish);
}
```

# The Quicksort Implementation

```
/*
 * Function: partition
 * -----
 * This function rearranges the elements of the vector so that the
 * small elements are grouped at the left end of the vector and the
 * large elements are grouped at the right end. The distinction
 * between small and large is made by comparing each element to the
 * pivot value, which is initially taken from vec[start]. When the
 * partitioning is done, the function returns a boundary index such
 * that vec[i] < pivot for all i < boundary, vec[i] == pivot
 * for i == boundary, and vec[i] >= pivot for all i > boundary.
 */

int partition(Vector<int> & vec, int start, int finish) {
    int pivot = vec[start];
    int lh = start + 1;
    int rh = finish;
    while (true) {
        while (lh < rh && vec[rh] >= pivot) rh--;
        while (lh < rh && vec[lh] < pivot) lh++;
        if (lh == rh) break;
        int tmp = vec[lh];
        vec[lh] = vec[rh];
        vec[rh] = tmp;
    }
    if (vec[lh] >= pivot) return start;
    vec[start] = vec[lh];
    vec[lh] = pivot;
    return lh;
}
```



## TUTORIAL

# The complexity of Quicksort

Worst case:  $O(N^2)$ , for a vector that is already sorted

Average case:  $O(N \log N)$ , for a random vector

- In practice, Quicksort is even much faster than Merge sort.
- [https://en.wikipedia.org/wiki/Sorting\\_algorithm](https://en.wikipedia.org/wiki/Sorting_algorithm)

Sorting complexity	Best	Average	Worst
Selection sort	$N^2$	$N^2$	$N^2$
Insertion sort	$N$	$N^2$	$N^2$
Bubble sort	$N$	$N^2$	$N^2$
Merge sort	$N \log N$	$N \log N$	$N \log N$
Quicksort	$N \log N$	$N \log N$	$N^2$

- There are many other factors to be considered when evaluating a sorting algorithm, or any other algorithms, which will be discussed in an advanced algorithm course.





# Other issues

- $O$  (bounded above) vs.  $\Omega$  (below) vs.  $\Theta$  (both)
  - E.g., the worst case of Quicksort is actually both  $O(N^2)$  and  $\Omega(N^2)$ , therefore  $\Theta(N^2)$  too.
  - An upper bound of the worst case is a bound on the running time of the algorithm on every input.
- Comparison sorting vs. other sorting
  - Worst case of any comparison sort is  $\Omega(N \log N)$ .
- Number of comparisons vs. number of swaps
- Time complexity vs. space complexity
- In-place (at most  $O(1)$  extra space) vs. not-in-place
- Recursive vs. non-recursive
- Stable vs. unstable
- Serial vs. parallel

*And all these factors affect each other.*

The End