# Lecture 27: Searching and Sorting 

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- Looking up a phone number, accessing a website and checking the definition of a word in a dictionary all involve searching large amounts of data.
- Searching algorithms all accomplish the same goal-finding an element that matches a given search key, if such an element does, in fact, exist.
- The major difference is the amount of effort they require to complete the search.
- One way to describe this effort is with Big O notation.
- For searching and sorting algorithms, this is particularly dependent on the number of data elements.
- In Chapter 7, we discussed the linear search algorithm, which is a simple and easy-toimplement searching algorithm.
- We'll now discuss the efficiency of the linear search algorithm as measured by Big O notation.
- Then, we'll introduce a searching algorithm that is relatively efficient but more complex to implement.


### 19.3. 1 Eificiency oi Linear Seapch

- Suppose an algorithm simply tests whether the first element of a vector is equal to the second element of the vector.
- If the vector has 10 elements, this algorithm requires only one comparison.
- If the vector has 1000 elements, the algorithm still requires only one comparison.
- In fact, the algorithm is independent of the number of vector elements.
- This algorithm is said to have a constant runtime, which is represented in Big O notation as $\mathrm{O}(1)$.
- An algorithm that is $O(1)$ does not necessarily require only one comparison.
- $\mathrm{O}(1)$ just means that the number of comparisons is constant-it does not grow as the size of the vector increases.
- An algorithm that tests whether the first element of a vector is equal to any of the next three elements will always require three comparisons, but in Big O notation it's still considered $\mathrm{O}(1)$.
- $O(1)$ is often pronounced "on the order of 1 " or more simply "order 1."
- An algorithm that tests whether the first element of a vector is equal to any of the other elements of the vector requires at most $n$ -1 comparisons, where n is the number of elements in the vector.
- If the vector has 10 elements, the algorithm requires up to nine comparisons.
- If the vector has 1000 elements, the algorithm requires up to 999 comparisons.
- As $n$ grows larger, the $n$ part of the expression "dominates," and subtracting one becomes inconsequential.
- Big O is designed to highlight these dominant terms and ignore terms that become unimportant as $n$ grows.


### 19.3.1 Efficiency oi Linear Search (conto

- An algorithm that requires a total of $n-1$ comparisons is said to be $\mathrm{O}(n)$.
- An $\mathrm{O}(n)$ algorithm is referred to as having a linear runtime.
- $\mathrm{O}(n)$ is often pronounced "on the order of $n$ " or more simply "order n."
- Now suppose you have an algorithm that tests whether any element of a vector is duplicated elsewhere in the vector.
- The first element must be compared with every other element in the vector.
- The second element must be compared with every other element except the first (it was already compared to the first).
- The third element must be compared with every other element except the first two.
- In the end, this algorithm will end up making $(n-1)+(n-2)+$ $\ldots+2+1$ or $n^{2} / 2-n / 2$ comparisons.
- As $n$ increases, the $n^{2}$ term dominates and the $n$ term becomes inconsequential.
- Again, Big O notation highlights the $n^{2}$ term, leaving $n^{2} / 2$.
- Big O is concerned with how an algorithm's runtime grows in relation to the number of items processed.
- Suppose an algorithm requires $n^{2}$ comparisons.
- With four elements, the algorithm will require 16 comparisons; with eight elements, 64 comparisons.
- With this algorithm, doubling the number of elements quadruples the number of comparisons.
- Consider a similar algorithm requiring $n^{2} / 2$ comparisons.
- With four elements, the algorithm will require eight comparisons; with eight elements, 32 comparisons.
- Again, doubling the number of elements quadruples the number of comparisons.
- Both of these algorithms grow as the square of $n$, so Big O ignores the constant, and both algorithms are considered to be $\mathrm{O}\left(n^{2}\right)$, which is referred to as quadratic runtime and pronounced "on the order of n -squared" or more simply "order n-squared."


### 19.3. 1 Eificiency oi Linear Seapch (conta)

- When n is small, $\mathrm{O}\left(n^{2}\right)$ algorithms will not noticeably affect performance.
- As $n$ grows, you'll start to notice the performance degradation.
- An $\mathrm{O}\left(n^{2}\right)$ algorithm running on a million-element vector would require a trillion "operations" (where each could actually require several machine instructions to execute).
- This could require a few hours to execute.
- A billion-element vector would require a quintillion operations, a number so large that the algorithm could take decades! Unfortunately, $\mathrm{O}\left(n^{2}\right)$ algorithms tend to be easy to write.
- In this chapter, you'll see algorithms with more favorable Big O measures.
- These efficient algorithms often take a bit more cleverness and effort to create, but their superior performance can be worth the extra effort, especially as $n$ gets large and as algorithms are compounded into larger programs.
- The linear search algorithm runs in $O(n)$ time.
- The worst case in this algorithm is that every element must be checked to determine whether the search key exists in the vector.
- If the size of the vector is doubled, the number of comparisons that the algorithm must perform is also doubled.
- Linear search can provide outstanding performance if the element matching the search key happens to be at or near the front of the vector.
- But we seek algorithms that perform well, on average, across all searches, including those where the element matching the search key is near the end of the vector.
- If a program needs to perform many searches on large vectors, it may be better to implement a different, more efficient algorithm, such as the binary search which we present in the next section.


## Performance Tip 19.1

Sometimes the simplest algorithms perform poorly. Their virtue is that they're easy to program, test and debug. Sometimes more complex algorithms are required to realize maximum performance.

## 19.3 .2 Binary Search

- The binary search algorithm is more efficient than the linear search algorithm, but it requires that the vector first be sorted.
- This is only worthwhile when the vector, once sorted, will be searched a great many times-or when the searching application has stringent performance requirements.
- The first iteration of this algorithm tests the middle element in the vector.
- If this matches the search key, the algorithm ends.


### 19.3.2 Binary Search (conti)

- Assuming the vector is sorted in ascending order, then if the search key is less than the middle element, the search key cannot match any element in the second half of the vector and the algorithm continues with only the first half of the vector (i.e., the first element up to, but not including, the middle element).
- If the search key is greater than the middle element, the search key cannot match any element in the first half of the vector and the algorithm continues with only the second half of the vector (i.e., the element after the middle element through the last element).
- Each iteration tests the middle value of the remaining portion of the vector.
- If the element does not match the search key, the algorithm eliminates half of the remaining elements.
- The algorithm ends either by finding an element that matches the search key or by reducing the subvector to zero size.

```
I // Fig 19.2: BinarySearch.h
// Class that contains a vector of random integers and a function
// that uses binary search to find an integer.
#include <vector>
using namespace std;
class BinarySearch
{
public:
    BinarySearch( int ); // constructor initializes vector
        int binarySearch( int ) const; // perform a binary search on vector
        void displayElements() const; // display vector elements
    private:
        int size; // vector size
        vector< int > data; // vector of ints
        void displaySubElements( int, int ) const; // display range of values
    }; // end class BinarySearch
```

Fig. 19.2 | BinarySearch class definition.

```
// Fig 19.3: BinarySearch.cpp
// BinarySearch class member-function definition.
#include <iostream>
#include <cstdlib> // prototypes for functions srand and rand
#include <ctime> // prototype for function time
#include <algorithm> // prototype for sort function
#include "BinarySearch.h" // class BinarySearch definition
using namespace std;
// constructor initializes vector with random ints and sorts the vector
BinarySearch::BinarySearch( int vectorSize )
{
    size = ( vectorSize > 0 ? vectorSize : 10 ); // validate vectorSize
    srand( time( 0 ) ); // seed using current time
    // fill vector with random ints in range 10-99
    for ( int i = 0; i < size; i++ )
            data.push_back( 10 + rand() % 90 ); // 10-99
        std::sort( data.begin(), data.end() ); // sort the data
} // end BinarySearch constructor
```

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Fig. 19.3 | BinarySearch class member-function definition. (Part I of 4.)

```
// perform a binary search on the data
int BinarySearch::binarySearch( int searchElement ) const
{
        int low = 0; // low end of the search area
        int high = size - 1; // high end of the search area
        int middle = ( low + high + 1 ) / 2; // middle element
        int location = -1; // return value; -1 if not found
        do // loop to search for element
        {
            // print remaining elements of vector to be searched
            displaySubElements( low, high );
        // output spaces for alignment
        for ( int i = 0; i < middle; i++ )
            cout << " ";
        cout << " * " << endl; // indicate current middle
        // if the element is found at the middle
        if ( searchElement == data[ middle ] )
            location = middle; // location is the current middle
        else if ( searchElement < data[ middle ] ) // middle is too high
        high = middle - 1; // eliminate the higher half
```

Fig. 19.3 | BinarySearch class member-function definition. (Part 2 of 4.)

```
        else // middle element is too low
            low = middle + 1; // eliminate the lower half
        middle = ( low + high + 1 ) / 2; // recalculate the middle
    } while ( ( low <= high ) && ( location == -1 ) );
    return location; // return location of search key
} // end function binarySearch
// display values in vector
void BinarySearch::displayElements() const
{
    displaySubElements( 0, size - 1 );
} // end function displayElements
// display certain values in vector
void BinarySearch::displaySubElements( int low, int high ) const
{
    for ( int i = 0; i < low; i++ ) // output spaces for alignment
        cout << " ";
    for ( int i = low; i <= high; i++ ) // output elements left in vector
        cout << data[ i ] << " ";
```

Fig. 19.3 | BinarySearch class member-function definition. (Part 3 of 4.)

71 cout << end1;
72 \} // end function displaySubElements
Fig. 19.3 | BinarySearch class member-function definition. (Part 4 of 4.)

### 19.3.2 Binary Seavch (conia)

- Line 29 initializes the location of the found element to -1-the value that will be returned if the search key is not found.
- Lines 31-51 loop until 1ow is greater than high (this occurs when the element is not found) or 1ocation does not equal -1 (indicating that the search key was found).
- Line 43 tests whether the value in the midd7e element is equal to searchElement.
- If so, line 44 assigns midd7e to location.
- Then the loop terminates and location is returned to the caller.
- Each iteration of the loop tests a single value (line 43) and eliminates half of the remaining values in the vector (line 46 or 48).

```
// Fig 19.4: Fig19_04.cpp
// BinarySearch test program.
#include <iostream>
#include "BinarySearch.h" // class BinarySearch definition
using namespace std;
    int main()
{
    int searchInt; // search key
    int position; // location of search key in vector
    // create vector and output it
    BinarySearch searchVector ( 15 );
    searchVector.displayElements();
    // get input from user
    cout << "\nP7ease enter an integer value (-1 to quit): ";
    cin >> searchInt; // read an int from user
    cout << endl;
    // repeatedly input an integer; -1 terminates the program
    while ( searchInt != -1 )
    {
```

Fig. 19.4 | BinarySearch test program. (Part I of 4.)

```
// use binary search to try to find integer
position = searchVector.binarySearch( searchInt );
// return value of -1 indicates integer was not found
if ( position == -1 )
        cout << "The integer " << searchInt << " was not found.\n";
else
            cout << "The integer " << searchInt
                << " was found in position " << position << ".\n";
                // get input from user
                cout << "\n\nP7ease enter an integer value (-1 to quit): ";
                cin >> searchInt; // read an int from user
                cout << endl;
            } // end while
} // end main
```

Fig. 19.4 | BinarySearch test program. (Part 2 of 4.)

```
26}313333847494967 73 74 82 89 90 91 95,
Please enter an integer value (-1 to quit): 38
26}33133\38474949 67 73 74 82 89 90 91 95,
26 31 33 384474949
The integer 38 was found in position 3.
Please enter an integer value (-1 to quit): 91
26}33133\384749494967 73 74 82 89 90 91 95,
    73 74 82 89 % 90 91 95
    90 91 95
```

The integer 91 was found in position 13.

Fig. 19.4 | BinarySearch test program. (Part 3 of 4.)

```
Please enter an integer value (-1 to quit): 25
26}34133\384749494967 73 74 82 89 90 91 95
26}313433847494
26 31 33
26
The integer 25 was not found.
Please enter an integer value (-1 to quit): -1
```

Fig. 19.4 | BinarySearch test program. (Part 4 of 4.)

- In the worst-case scenario, searching a sorted vector of 1023 elements will take only 10 comparisons when using a binary search.
- Repeatedly dividing 1023 by 2 (because, after each comparison, we can eliminate from consideration half of the remaining vector) and rounding down (because we also remove the middle element) yields the values $511,255,127$, $63,31,15,7,3,1$ and 0.
- The number $1023\left(2^{10}-1\right)$ is divided by 2 only 10 times to get the value 0 , which indicates that there are no more elements to test.


### 19.3.2 Binary Search (conto)

- Dividing by 2 is equivalent to one comparison in the binary search algorithm.
- Thus, a vector of 1,048,575 $\left(2^{20}-1\right)$ elements takes a maximum of 20 comparisons to find the key, and a vector of about one billion elements takes a maximum of 30 comparisons to find the key.
- This is a tremendous improvement in performance over the linear search.
- For a one-billion-element vector, this is a difference between an average of 500 million comparisons for the linear search and a maximum of only 30 comparisons for the binary search! The maximum number of comparisons needed for the binary search of any sorted vector is the exponent of the first power of 2 greater than the number of elements in the vector, which is represented as $\log _{2} n$.


### 19.3.2 圂inary Seapch (cont.)

- All logarithms grow at roughly the same rate, so in Big O notation the base can be omitted.
- This results in a Big O of $\mathrm{O}(\log n)$ for a binary search, which is also known as logarithmic runtime and pronounced "on the order of $\log$ n" or more simply "order $\log n$."


### 19.4 Soriting Algorithns

- Sorting data (i.e., placing the data into some particular order, such as ascending or descending) is one of the most important computing applications.
- The choice of algorithm affects only the runtime and memory use of the program.
- The next section examines the efficiency of selection sort and insertion sort algorithms using Big O notation.
- The last algorithm-merge sort, which we introduce in this chapter-is much faster but is more difficult to implement.


### 19.4. 1 Eificiency of Selection Sort

- Selection sort is an easy-to-implement, but inefficient, sorting algorithm.
- The first iteration of the algorithm selects the smallest element in the vector and swaps it with the first element.
- The second iteration selects the second-smallest element (which is the smallest element of the remaining elements) and swaps it with the second element.
- The algorithm continues until the last iteration selects the second-largest element and swaps it with the second-to-last element, leaving the largest element in the last index.
- After the $i^{\text {th }}$ iteration, the smallest $i$ elements of the vector will be sorted into increasing order in the first $i$ elements of the vector.


### 19.4. 1 Eiticiency oi selection sori (conid)

- The selection sort algorithm iterates $n-1$ times, each time swapping the smallest remaining element into its sorted position.
- Locating the smallest remaining element requires $n-$ 1 comparisons during the first iteration, $n-2$ during the second iteration, then $n-3, \ldots, 3,2,1$.
- This results in a total of $n(n-1) / 2$ or $\left(n^{2}-n\right) / 2$ comparisons.
- In Big O notation, smaller terms drop out and constants are ignored, leaving a final Big O of $O\left(n^{2}\right)$.


## Insertion sort is another simple, but inefficient, sorting algorithm.

- The algorithm's first iteration takes the second element in the vector and, if it's less than the first element, swaps it with the first element.
- The second iteration looks at the third element and inserts it into the correct position with respect to the first two elements, so all three elements are in order.
- At the $i^{t h}$ iteration of this algorithm, the first $i$ elements in the original vector will be sorted.
- Insertion sort iterates $n-1$ times, inserting an element into the appropriate position in the elements sorted so far.


### 19.4 2 Eiriciency oi Inseriton sorit (contu)

- For each iteration, determining where to insert the element can require comparing the element to each of the preceding elements- $n-1$ comparisons in the worst case.
- Each individual repetition statement runs in $O(n)$ time.
- For determining Big O notation, nested statements mean that you must multiply the number of comparisons.
- For each iteration of an outer loop, there will be a certain number of iterations of the inner loop.
- In this algorithm, for each $O(n)$ iteration of the outer loop, there will be $O(n)$ iterations of the inner loop, resulting in a Big O of $O(n * n)$ or $O\left(n^{2}\right)$.


### 19.4.3 Mergo Sort (A Recursive <br> Implementation)

- Merge sort is an efficient sorting algorithm but is conceptually more complex than selection sort and insertion sort.
- The merge sort algorithm sorts a vector by splitting it into two equal-sized subvectors, sorting each subvector then merging them into one larger vector.
- Merge sort performs the merge by looking at the first element in each vector, which is also the smallest element in the vector.
- Merge sort takes the smallest of these and places it in the first element of the larger, sorted vector.
- If there are still elements in the subvector, merge sort looks at the second element in that subvector (which is now the smallest element remaining) and compares it to the first element in the other subvector.
- Merge sort continues this process until the larger vector is filled.


### 19.4.3 Merge Sort (A Recursive Implementation) (conto)

- The implementation of merge sort in this example is recursive.
- The base case is a vector with one element.
- A one-element vector is, of course, sorted, so merge sort immediately returns when it's called with a one-element vector.
- The recursion step splits a vector of two or more elements into two equal-sized subvectors, recursively sorts each subvector, then merges them into one larger, sorted vector.
- If there is an odd number of elements, one subvector is one element larger than the other.


### 19.4.3 Merge Sort (A Recursive <br> Implementation) (conto)

- Figure 19.5 defines class MergeSort, and lines 22-25 of Fig. 19.6 define the sort function.
- Line 24 calls function sortSubvector with 0 and size
-1 as the arguments.
- These arguments correspond to the beginning and ending indices of the vector to be sorted, causing sortSubVector to operate on the entire vector.
- Function sortSubVector is defined in lines 28-52.
- Line 31 tests the base case.
- If the size of the vector is 0 , the vector is already sorted, so the function simply returns immediately.
- If the size of the vector is greater than or equal to 1 , the function splits the vector in two, recursively calls function sortSubVector to sort the two subvectors, then merges them.


### 19.4.3 Merge Sort (A Recursive Innplennentation) (conty)

- Line 46 recursively calls function sortSubVector on the first half of the vector, and line 47 recursively calls function sortSubVector on the second half of the vector.
- When these two function calls return, each half of the vector has been sorted.
- Line 50 calls function merge (lines 55-99) on the two halves of the vector to combine the two sorted vectors into one larger sorted vector.

```
// Fig 19.5: MergeSort.h
// Class that creates a vector filled with random integers.
// Provides a function to sort the vector with merge sort.
#include <vector>
using namespace std;
// MergeSort class definition
class MergeSort
{
public:
    MergeSort( int ); // constructor initializes vector
    void sort(); // sort vector using merge sort
    void displayElements() const; // display vector elements
    private:
        int size; // vector size
        vector< int > data; // vector of ints
        void sortSubVector( int, int ); // sort subvector
        void merge( int, int, int, int ); // merge two sorted vectors
        void displaySubVector( int, int ) const; // display subvector
    }; // end class SelectionSort
```

Fig. I9.5 $\mid$ MergeSort class definition.

```
// Fig 19.6: MergeSort.cpp
// Class MergeSort member-function definition.
#include <iostream>
#include <vector>
#include <cstdlib> // prototypes for functions srand and rand
#include <ctime> // prototype for function time
#include "MergeSort.h" // class MergeSort definition
using namespace std;
// constructor fil1 vector with random integers
MergeSort::MergeSort( int vectorSize )
{
        size = ( vectorSize > 0 ? vectorSize : 10 ); // validate vectorSize
        srand( time( 0 ) ); // seed random number generator using current time
        // fill vector with random ints in range 10-99
        for ( int i = 0; i < size; i++ )
            data.push_back( 10 + rand() % 90 );
} // end MergeSort constructor
    // split vector, sort subvectors and merge subvectors into sorted vector
    void MergeSort::sort()
{
```

Fig. 19.6 | MergeSort class member-function definition. (Part I of 5.)

```
    sortSubVector( 0, size - 1 ); // recursively sort entire vector
} // end function sort
// recursive function to sort subvectors
void MergeSort::sortSubVector( int low, int high )
{
    // test base case; size of vector equals 1
    if ( ( high - low ) >= 1 ) // if not base case
    {
        int middle1 = ( low + high ) / 2; // calculate middle of vector
        int middle2 = middle1 + 1; // calculate next element over
        // output sp7it step
        cout << "split: ";
        displaySubVector( low, high );
        cout << endl << " ";
        displaySubVector( low, middle1 );
        cout << endl << " ";
        displaySubVector( middle2, high );
        cout << endl << endl;
        // split vector in half; sort each half (recursive calls)
        sortSubVector( low, middle1 ); // first half of vector
        sortSubVector( middle2, high ); // second half of vector
```

Fig. 19.6 | MergeSort class member-function definition. (Part 2 of 5.)

```
4 8
4 9
5 0
5 1
5 2
5 3
5 4
5 5
5 6
5 7
5 8
5 9
6 0
```

        // merge two sorted vectors after split calls return
    ```
        // merge two sorted vectors after split calls return
        merge( low, middle1, middle2, high );
        merge( low, middle1, middle2, high );
        } // end if
        } // end if
} // end function sortSubVector
} // end function sortSubVector
// merge two sorted subvectors into one sorted subvector
// merge two sorted subvectors into one sorted subvector
void MergeSort::merge( int left, int middle1, int middle2, int right )
void MergeSort::merge( int left, int middle1, int middle2, int right )
{
{
    int leftIndex = 1eft; // index into left subvector
    int leftIndex = 1eft; // index into left subvector
    int rightIndex = middle2; // index into right subvector
    int rightIndex = middle2; // index into right subvector
    int combinedIndex = left; // index into temporary working vector
    int combinedIndex = left; // index into temporary working vector
    vector< int > combined( size ); // working vector
    vector< int > combined( size ); // working vector
    // output two subvectors before merging
    // output two subvectors before merging
    cout << "merge: ";
    cout << "merge: ";
    displaySubVector( left, middle1 );
    displaySubVector( left, middle1 );
    cout << endl << " ";
    cout << endl << " ";
    disp1aySubVector( middle2, right );
    disp1aySubVector( middle2, right );
    cout << endl;
    cout << endl;
    // merge vectors until reaching end of either
    // merge vectors until reaching end of either
    while ( leftIndex <= middle1 && rightIndex <= right )
    while ( leftIndex <= middle1 && rightIndex <= right )
    {
```

    {
    ```

Fig. 19.6 | MergeSort class member-function definition. (Part 3 of 5.)
```

7 2
7 3
7 4
7 5
7 6
7 7
78
7 9
80
8I
82
83
84
85
86
87
88
89
90
9 1
92
93

```
```

        // place smaller of two current elements into result
    ```
        // place smaller of two current elements into result
        // and move to next space in vector
        // and move to next space in vector
        if ( data[ leftIndex ] <= data[ rightIndex ] )
        if ( data[ leftIndex ] <= data[ rightIndex ] )
            combined[ combinedIndex++ ] = data[ leftIndex++ ];
            combined[ combinedIndex++ ] = data[ leftIndex++ ];
        else
        else
        combined[ combinedIndex++ ] = data[ rightIndex++ ];
        combined[ combinedIndex++ ] = data[ rightIndex++ ];
} // end while
} // end while
if ( leftIndex == middle2 ) // if at end of left vector
if ( leftIndex == middle2 ) // if at end of left vector
{
{
        while ( rightIndex <= right ) // copy in rest of right vector
        while ( rightIndex <= right ) // copy in rest of right vector
            combined[ combinedIndex++ ] = data[ rightIndex++ ];
            combined[ combinedIndex++ ] = data[ rightIndex++ ];
} // end if
} // end if
else // at end of right vector
else // at end of right vector
{
{
    while ( leftIndex <= middle1 ) // copy in rest of left vector
    while ( leftIndex <= middle1 ) // copy in rest of left vector
            combined[ combinedIndex++ ] = data[ leftIndex++ ];
            combined[ combinedIndex++ ] = data[ leftIndex++ ];
} // end else
} // end else
// copy values back into original vector
// copy values back into original vector
for ( int i = left; i <= right; i++ )
for ( int i = left; i <= right; i++ )
    data[ i ] = combined[ i ];
```

    data[ i ] = combined[ i ];
    ```
94

Fig. 19.6 | MergeSort class member-function definition. (Part 4 of 5.)
```

95 // output merged vector
96 cout << " ";
97 displaySubVector( left, right );
98 cout << endl << endl;
99 } // end function merge
100
IOI // display elements in vector
102 void MergeSort::displayElements() const
103 {
104 displaySubVector( 0, size - 1 );
I05 } // end function displayElements
l06
I07 // display certain values in vector
I08 void MergeSort::displaySubVector( int low, int high ) const
109 {
// output spaces for alignment
for ( int i = 0; i < low; i++ )
cout << ";
// output elements left in vector
for ( int i = low; i <= high; i++ )
cout << " " << data[ i ];
} // end function displaySubVector

```

Fig. 19.6 | MergeSort class member-function definition. (Part 5 of 5.)

\subsection*{19.4.3 Merge Sort (A Recursive Implementation) (conto)}
- Lines 70-78 in function merge loop until the program reaches the end of either subvector.
- Line 74 tests which element at the beginning of the vectors is smaller.
- If the element in the left vector is smaller, line 75 places it in position in the combined vector.
- If the element in the right vector is smaller, line 77 places it in position in the combined vector.
- When the whi 1 e loop has completed (line 78), one entire subvector is placed in the combined vector, but the other subvector still contains data.
- Line 80 tests whether the left vector has reached the end.
- If so, lines \(82-83\) fill the combined vector with the elements of the right vector.

\subsection*{19.4.3 Merge Sort (A Recursive \\ In plennentation) (conty}
- If the left vector has not reached the end, then the right vector must have reached the end, and lines 8788 fill the combined vector with the elements of the left vector.
- Finally, lines 92-93 copy the combined vector into the original vector.
- Figure 19.7 creates and uses a MergeSort object.
- The output from this program displays the splits and merges performed by merge sort, showing the progress of the sort at each step of the algorithm.
```

// Fig 19.7: Fig19_07.cpp
// MergeSort test program.
\#include <iostream>
\#include "MergeSort.h" // class MergeSort definition
using namespace std;
int main()
{
// create object to perform merge sort
MergeSort sortVector( 10 );
cout << "Unsorted vector:" << endl;
sortVector.displayElements(); // print unsorted vector
cout << endl << endl;
sortVector.sort(); // sort vector
cout << "Sorted vector:" << endl;
sortVector.displayElements(); // print sorted vector
cout << endl;
} // end main

```

Fig. \(19.7 \left\lvert\, \begin{array}{ll}\text { | }\end{array}\right.\)
```

Unsorted vector:
30 47 22 67 79 18 60 78 26 54
split: 30 47 22 67 79 18 60 78 26 54

```

```

split: }304722677
3047 22
6779
split: 3047 22
3047
22
split: 3047
30
4 7
merge: 30
3047

```

Fig. 19.7 \(\mid\) MergeSort test program. (Part 2 of 5.)
```

merge: 3047
22 3047
split:
6 7 7 9
6 7
7 9
merge: 67 79
6 7 7 9
merge: 22 30 47
22 3047 67 79
split:
18 6078 26 54
18 60 78
26 54
split:
18 60 78
78

```

Fig. \(19.7 \left\lvert\, \begin{array}{ll}\text { | }\end{array}\right.\) MergeSort test program. (Part 3 of 5.)


Fig. \(19.7 \left\lvert\, \begin{array}{ll}\text { | }\end{array}\right.\) MergeSort test program. (Part 4 of 5.)
```

merge: 22 30 47 67 79
18}222\quad26 30 47 54 60 67 78 79

```
Sorted vector:
    \(\begin{array}{llllllll}18 & 22 & 26 & 30 & 47 & 54 & 60 & 67 \\ 78 & 79\end{array}\)

Fig. \(19.7 \mid\) MergeSort test program. (Part 5 of 5.)

\subsection*{19.4.3 Merge Sort (A Recursive Implementation) (conto)}
- Merge sort is a far more efficient algorithm than either insertion sort or selection sort.
- Consider the first (nonrecursive) call to function sortSubVector (line 24).
- This results in two recursive calls to function sortSubVector with subvectors each approximately half the size of the original vector, and a single call to function merge.
- This call to function merge requires, at worst, \(n-1\) comparisons to fill the original vector, which is \(\mathrm{O}(\mathrm{n})\).
- The two calls to function sortSubVector result in four more recursive calls to function sortSubVector-each with a subvector approximately one-quarter the size of the original vector-and two calls to function merge.
- These two calls to function merge each require, at worst, \(n / 2-1\) comparisons, for a total number of comparisons of \(O(n)\).

\subsection*{19.4.3 Merge Sort (A Recursive Implementation) (conto)}
- This process continues, each call to sortSubVector generating two additional calls to sortSubvector and a call to merge, until the algorithm has split the vector into one-element subvectors.
- At each level, \(O(n)\) comparisons are required to merge the subvectors.
- Each level splits the size of the vectors in half, so doubling the size of the vector requires one more level.
- Quadrupling the size of the vector requires two more levels.
- This pattern is logarithmic and results in \(\log _{2} n\) levels.
- This results in a total efficiency of \(\mathrm{O}(n \log n)\).

\subsection*{19.4.3 Merge Sort (A Recursive Implementation) (conto)}
- Figure 19.8 summarizes the searching and sorting algorithms we cover in this book and lists the Big O for each.
- Figure 19.9 lists the Big O categories we've covered in this chapter along with a number of values for \(n\) to highlight the differences in the growth rates.
\begin{tabular}{lll|}
\hline Algorithm & Location & Big O \\
Searching Algorithms & & \\
Linear search & Section 7.7 & \(O(n)\) \\
Binary search & Section 19.2 .2 & \(O(\log n)\) \\
Recursive linear search & Exercise 19.8 & \(O(n)\) \\
Recursive binary search & Exercise 19.9 & \(O(\log n)\) \\
Sorting Algorithms & & \\
Insertion sort & Section 7.8 & \(O\left(n^{2}\right)\) \\
Selection sort & Section 8.6 & \(O\left(n^{2}\right)\) \\
Merge sort & Section 19.3 .3 & \(O(n \log n)\) \\
Bubble sort & Exercises 19.5 and 19.6 & \(O\left(n^{2}\right)\) \\
Quicksort & Exercise 19.10 & Worst case: \(O\left(n^{2}\right)\) \\
& & Average case: \(O(n \log n)\)
\end{tabular}

Fig. 19.8 | Searching and sorting algorithms with Big O values.
\begin{tabular}{|llllll|}
\hline & \begin{tabular}{l} 
Approximate \\
decimal value
\end{tabular} & \(O(\log n)\) & \(O(n)\) & \(O(n \log n)\) & \(O\left(n^{2}\right)\) \\
\(n\) & 1000 & 10 & \(2^{10}\) & \(10 \cdot 2^{10}\) & \(2^{20}\) \\
\(2^{10}\) & \(1,000,000\) & 20 & \(2^{20}\) & \(20 \cdot 2^{20}\) & \(2^{40}\) \\
\(2^{20}\) & \(1,000,000,000\) & 30 & \(2^{30}\) & \(30 \cdot 2^{30}\) & \(2^{60}\) \\
\(2^{30}\) & & & & \\
\hline
\end{tabular}

Fig. 19.9 | Approximate number of comparisons for common Big O notations.

\section*{Questions}
```

