

This topic teaches the copy algorithm by demonstrating the usage including copy_backward, merge, unique and reverse and explaining the algorithms

• Figure 16.8 demonstrates algorithms copy_backward, merge, unique and reverse.

copy backward(a1.cbegin(), a1.cend(), results.end());

- Uses the copy backward algorithm to copy elements in the range from al.cbegin() up to, but *not* including, a1.cend()
 - Placing the elements in results by starting from the element before results.end()
 - Working toward the beginning of the array
- Returns an iterator positioned at the *last* element copied into the results (i.e., the beginning of results, because of the backward copy)
- The elements are placed in results in the same order as a1

- copy_backward can manipulate overlapping ranges of elements in a container as long as the first element to copy is not in the destination range of elementskward through a sequence, respectively
- One difference between copy_backward and copy is that
 - The iterator returned from copy is positioned after the last element copied
 - The one returned from copy_backward is positioned at the last element copied (i.e., the first element in the sequence)
- copy_backward *can* manipulate *overlapping* ranges of elements in a container as long as the first element to copy is *not* in the destination range of elements



- In addition to the copy and copy_backward algorithms
 - -C++11 now includes the move and move_backward algorithms

```
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```

```
merge(a1.cbegin(), a1.cend(),
      a2.cbegin(), a2.cend(),
      results2.begin());
```

- Uses the merge algorithm to combine two *sorted ascending* sequences of values into a third sorted ascending sequence
- Requires five iterator arguments
- The first four must be at least *input iterators*
- The last must be at least an *output iterator*
- The first two arguments specify the range of elements in the first sorted sequence (a1), the second two arguments specify the range of elements in the second sorted sequence (a2) and the last argument specifies the starting location in the third sequence (results2) where the elements will be merged
- A second version of this algorithm takes as its sixth argument a *binary* predicate function that specifies the sorting order

```
array< int, SIZE + SIZE > results2;
```

- Creates the array results 2 with the number of elements in a 1 and a2 (SIZE)
- Using the merge algorithm requires that the sequence where the results are stored be at least the size of the sequences being merged
- If you do not want to allocate the number of elements for the resulting sequence before the merge operation
 - You can use the following statements:

```
vector< int > results2;
merge(a1.begin(), a1.end(),
      a2.begin(), a2.end(),
      back inserter( results2 ) );
```

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- The argument back_inserter(results2) uses function template back inserter (header <iterator>) for the vector results2
- A back inserter calls the container's default push_back function to insert an element at the end of the container
- If an element is inserted into a container that has no more space available, the container grows in size which is why we used a vector in the preceding statements, because arrays are fixed size
- Thus, the number of elements in the container does *not* have to be known in advance

- There are two other inserters
 - -front inserter
 - Uses push_front to insert an element at the *beginning* of a container specified as its argument)
 - -Inserter
 - Uses insert to insert an element *at* the iterator supplied as its second argument in the container supplied as its first argument

auto endLocation = unique(results2.begin(), results2.end());

- Uses the unique algorithm on the sorted sequence of elements in the range from results2.begin() up to, but *not* including, results2.end()
- After this algorithm is applied to a sorted sequence with duplicate values, only a single copy of each value remains in the sequence
- Takes two arguments that must be at least forward iterators



- The algorithm returns an iterator positioned *after* the last element in the sequence of unique values
- The values of all elements in the container after the last unique value are *undefined*
- A second version of this algorithm takes as a third argument a binary predicate function specifying how to compare two elements for *equality*

reverse(a1.begin(), a1.end());

- Uses the reverse algorithm to reverse all the elements in the range from al.begin() up to, but not including, al.end()
- Takes two arguments that must be at least bidirectional iterators



copy_if

- Copies each element from a range if the *unary predicate* function in its fourth argument returns **true** for that element
- The iterators supplied as the first two arguments must be *input* iterators
- The iterator supplied as the third argument must be an *output iterator* so that the element being copied can be assigned to the copy location
- Returns an iterator positioned after the *last* element copied

copy_n

- Copies the number of elements specified by its second argument from the location specified by its first argument (an input iterator)
- The elements are output to the location specified by its third argument (an output iterator)

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