#### Divide-and-Conquer

Divide-and-conquer is a general algorithm design paradigm:

- Divide: divide the input data S in two disjoint subsets  $S_1$  and  $S_2$
- Recur: solve the subproblems associated with  $S_1$  and  $S_2$ 
  - the base case for the recursion are subproblems of size 0 or 1
- Conquer: combine the solutions for  $S_1$  and  $S_2$  into a solution for S

Merge-sort is a sorting algorithm based on the divide-and-conquer paradigm

- Like heap-sort
  - Uses a comparator
  - Has  $O(n \log n)$  running time
- Unlike heap-sort
  - Does not use an auxiliary priority queue
  - Accesses data in a sequential manner (suitable to sort data on a disk)

#### Merge Sort

Merge-sort on an input sequence S with n elements consists of three steps:

- Divide: partition S into two sequences  $S_1$  and  $S_2$  of about n/2 elements each
- Recur: recursively sort  $S_1$  and  $S_2$
- Conquer: merge  $S_1$  and  $S_2$  into a unique sorted sequence

```
Algorithm mergeSort(S, C)
Input sequence S with n elements, comparator C
Output sequence S sorted according to C
if S.size() > 1
(S_1, S_2) \leftarrow partition(S, n/2)
mergeSort(S_1, C)
mergeSort(S_2, C)
S \leftarrow merge(S_1, S_2)
```

#### Merging two sorted sequences

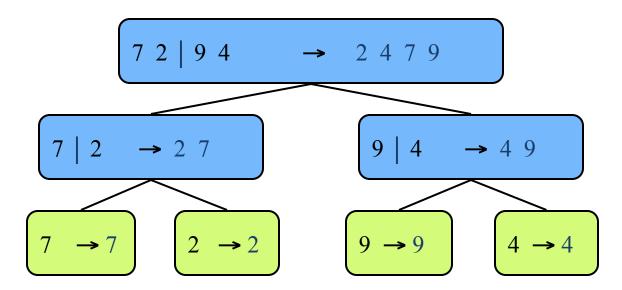
- The conquer step of merge-sort consists of merging two sorted sequences A and B into a sorted sequence S containing the union of the elements of A and B
- Merging two sorted sequences, each with n/2, takes O(n) time

```
Algorithm merge(A, B)
   Input sequences A and B with n/2 elements each
   Output sorted sequence of A \cup B
   S \leftarrow empty sequence
   while \neg A.isEmpty() \land \neg B.isEmpty()
       if A.first().element() < B.first().element()
           S.insertLast(A.remove(A.first()))
       else
           S.insertLast(B.remove(B.first()))
   while \neg A.isEmpty()
       S.insertLast(A.remove(A.first()))
   while \neg B.isEmpty()
       S.insertLast(B.remove(B.first()))
   return S
```

#### Merge-Sort Tree

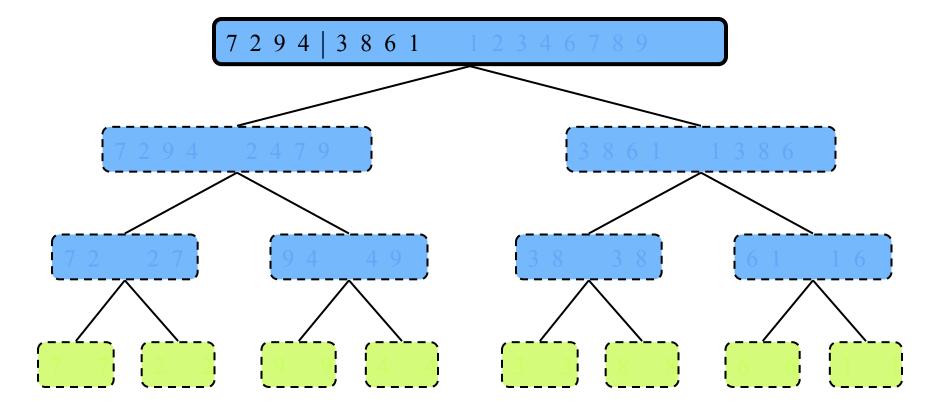
An execution of merge-sort is depicted by a binary tree

- each node represents a recursive call of merge-sort and stores
  - unsorted sequence before the execution and its partition
  - sorted sequence at the end of the execution
- the root is the initial call
- the leaves are calls on subsequences of size 1

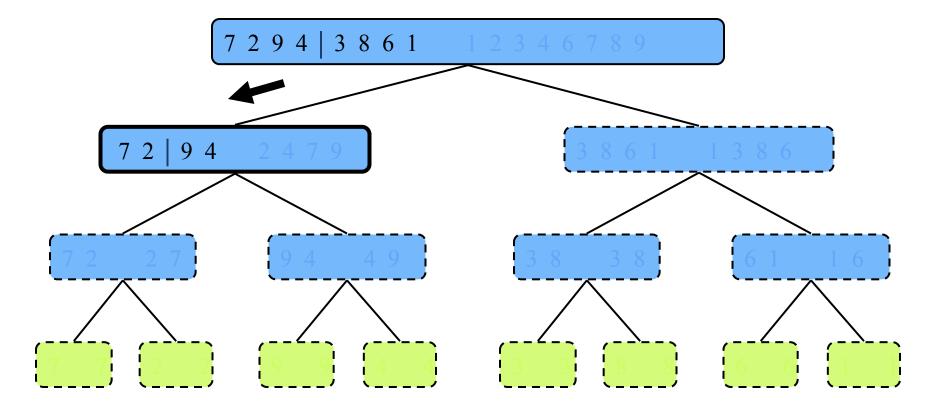


# **Execution Example**

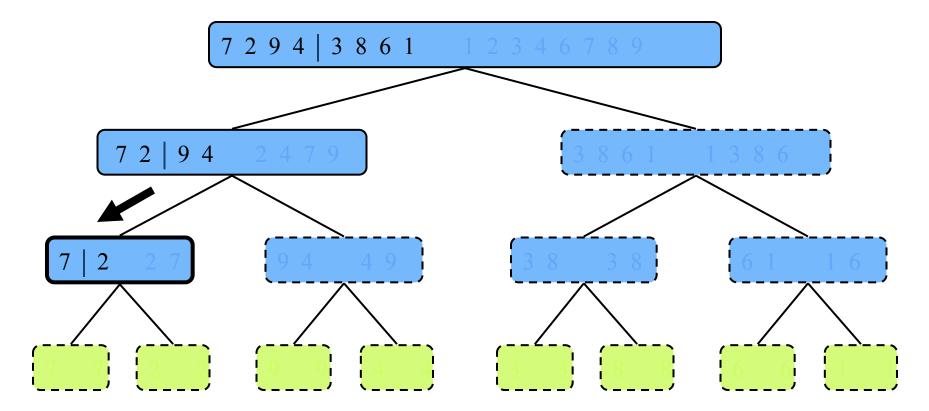
• Partition



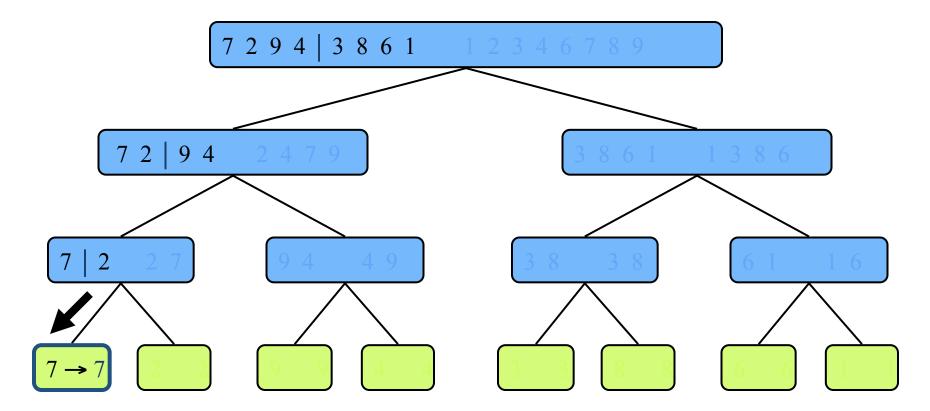
• Recursive call, partition



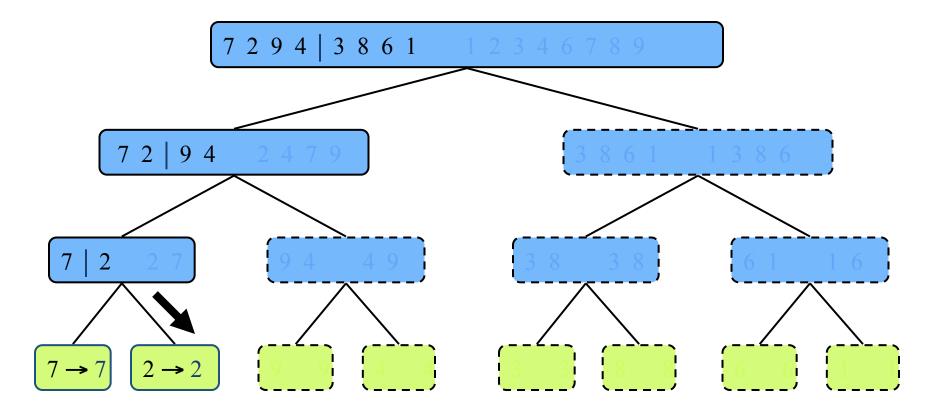
• Recursive call, partition



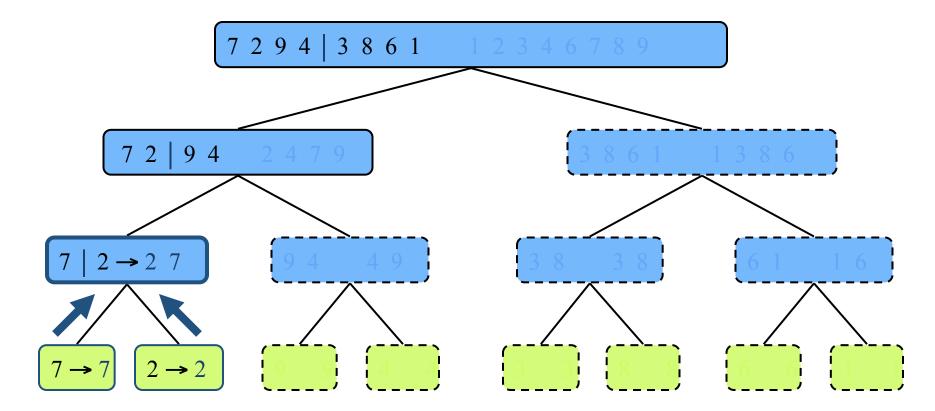
• Recursive call, base case



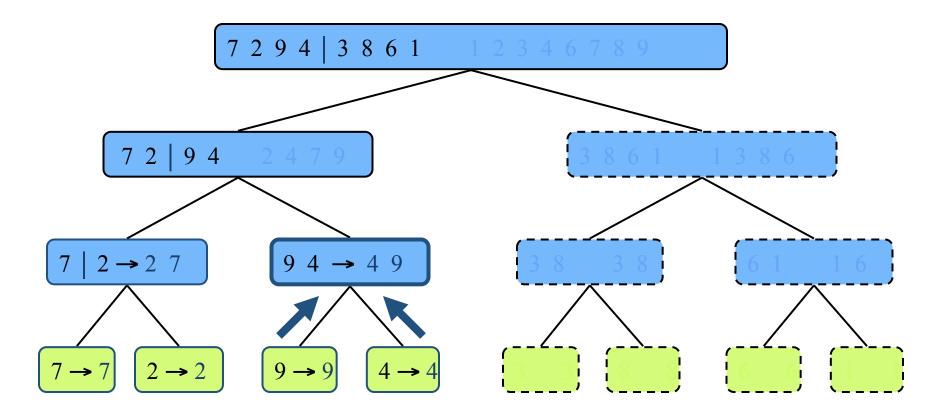
• Recursive call, base case



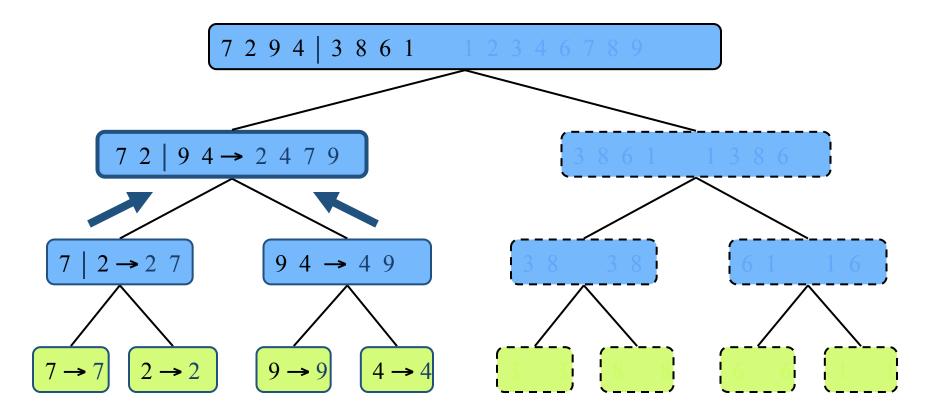
Merge



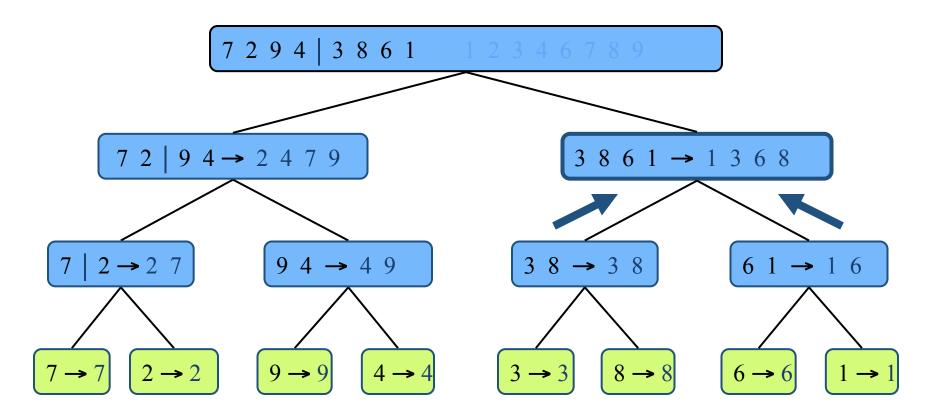
• Recursive call, ..., base case, merge



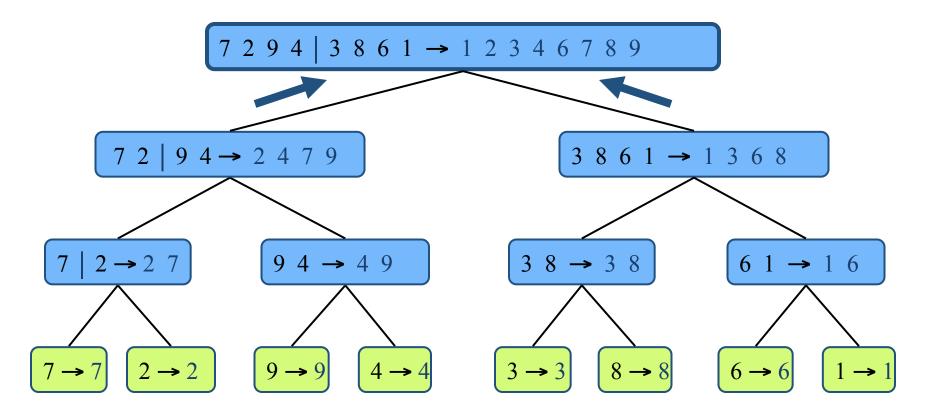
Merge



• Recursive call, ..., merge, merge



Merge



### Analysis of Merge-Sort

- The height h of the merge-sort tree is  $O(\log n)$ 
  - at each recursive call we divide the sequence in half
- The overall amount or work done at the nodes of depth i is O(n)
  - we partition and merge  $2^i$  sequences of size  $n/2^i$
  - we make  $2^{i+1}$  recursive calls
- Thus, the total running time of merge-sort is  $O(n \log n)$

#### 

### Comparing sorting algorithms

Consider the following when evaluating a sorting algorithm:

- Time complexity
- Space complexity
  - An in-place algorithm requires only n + O(1) space, using the already given space for the n elements and O(1) additional space
- Stability
  - A sorting algorithm is stable if it preserves the original relative ordering of elements with equal value
  - Ex: Unsorted sequence ( $\mathbf{B}$ ,  $\mathbf{b}$ ,  $\mathbf{a}$ ,  $\mathbf{c}$ ). Suppose  $\mathbf{B} = \mathbf{b}$  and  $\mathbf{a} < \mathbf{b} < \mathbf{c}$ .
    - Stable sorted: (a, **B**, b, c)
    - Unstable sorted: (a, b, B, c)
  - Necessary if we want to sort repeatedly by different attributes (i.e., sort by first name, then sort again by last name)

## Summary of Sorting Algorithms

Algorithm	Time	Notes
selection-sort	$O(n^2)$	<ul><li>♦ in-place</li><li>♦ not stable</li><li>♦ for small data sets (&lt; 1K)</li></ul>
insertion-sort	$O(n^2)$	<ul><li>♦ in-place</li><li>♦ stable</li><li>♦ for small data sets (&lt; 1K)</li></ul>
heap-sort	$O(n \log n)$	<ul> <li>in-place</li> <li>not stable</li> <li>for large data sets (1K — 1M)</li> </ul>
merge-sort	$O(n \log n)$	<ul> <li>not in-place</li> <li>stable</li> <li>sequential data access</li> <li>for huge data sets (&gt; 1M)</li> </ul>

#### Other

• You are given a query point *p* and a set *S* of *n* other points in two dimensional space. Find *k* points out of the *n* points which are nearest to *p*.