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RŪŠIAVIMO ALGORITMŲ ANIMACIJA:

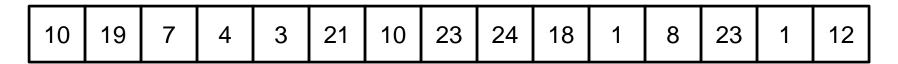
https://www.bluffton.edu/~nesterd/java/SortingDemo.html

Outline

- Motivation
- Quadratic O(n²) Sorting
 - Selection Sort
 - Insertion Sort
- "Linearithmic" O(n log n) Sorting
 - Merge Sort
 - Quick Sort
- The Master Theorem
- Linear Sorting
 - Radix Sort

Motivation

- Problem:
 - Turn this:



• Into this:

As efficiently as possible!

Sorting Algorithms

- There are many ways to sort arrays:
 - Iterative vs. recursive
 - In-place vs. not-in-place
 - An in-place algorithm transforms an input data structure with a small (constant) amount of extra storage space
 - In the context of sorting, this means that the input array is overwritten by the output as the algorithm executes instead of introducing a new array
 - Comparison-based vs. non-comparative
 - Most sorting algorithms we'll analyze are comparisonbased--that is, the sort is produced by comparing elements with each other--but some (like Radix Sort, which we'll see later) are not

"In-placeness"

Consider this function to reverse an array:

```
function reverse(A):
    n = A.length
    B = array of length n
    for i = 0 to n - 1:
        B[n-1-i] = A[i]
    return B
```

Not in-place!

Now consider this version:

```
function reverse(A):
    n = A.length
    for i = 0 to n/2:
        temp = A[i]
        A[i] = A[n-1-i]
        A[n-1-i] = temp
    // return statement not needed!
```

In-place!

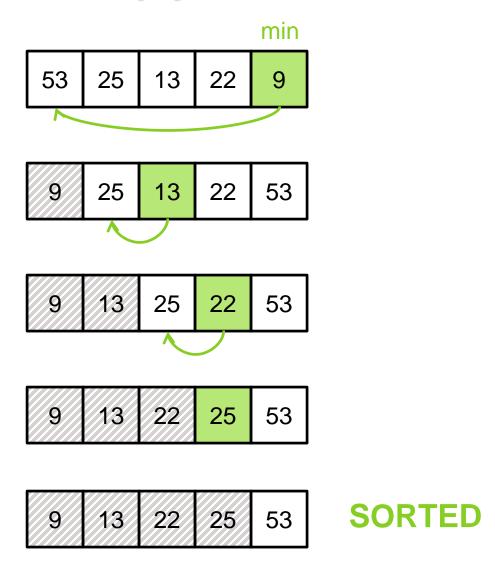
Advantages of in place solutions?

- They are sometimes more difficult to write, but take up much less memory
- Tradeoff between space efficiency and simplicity of the algorithm
- More difficult to write in-place solutions for recursive algorithms

Selection Sort

- Usually iterative and in-place
- Divides input array into two logical parts
 - elements already sorted
 - elements that still need to be sorted
- Selects the smallest element of the array and places it at index 0, then selects the second smallest and places it in index 1, then the third smallest in index 2, etc..
- Advantages:
 - Very simple
 - Memory efficient: in-place means swapping elements within same array
- Disadvantages:
 - Slow: runs in quadratic O(n²) time.

Selection Sort (2)



Selection Sort (3)

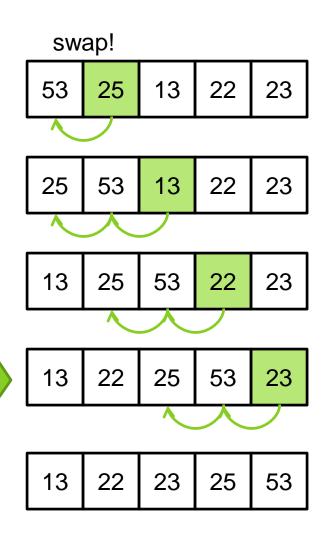
```
function selection_sort(A):
    // Input: Unsorted List
    // Output: Sorted List
    n = A.length
    for i = 0 to n-2:
        min = argmin(A[i:n-1])
        swap A[i] with A[min]
```

Insertion Sort

- Usually iterative and in-place
- Arranges items one at a time by comparing each element with every element before it and inserting it into the correct position
- Advantages:
 - Works particularly well if the list is already partially sorted (we'll see why)
 - Memory efficient: in-place means swapping elements within same array
- Disadvantages:
 - Slow: runs in **quadratic** O(n²) time.

Insertion Sort (2)

Note that since 23 > 22, we don't need to check any further, since that part of the array is already sorted. This demonstrates that insertion sort will cruise through an already sorted array in linear time!



Insertion Sort (3)

Divide-and-Conquer

- Divide-and-conquer is a general algorithm design paradigm:
 - Divide: divide the input data S into disjoint subsets S₁, S₂, ..., S_k
 - Recur: solve the subproblems associated with S₁, S₂, ..., S_k
 - Conquer: combine the solutions for S₁, S₂, ..., S_k into a solution for S
- The base case for the recursion is generally subproblems of size 0 or 1

Merge Sort

- Merge sort is a sorting algorithm based on the divide-and-conquer paradigm
- Like the quadratic sorts, merge sort is comparative
- Unlike the quadratic sorts, merge sort is recursive and runs in linearithmic, O(n log n), time!
 - Let's see why...

Merge Sort (2)

- Merge sort on an input sequence S with n elements consists of three steps:
 - Divide: partition S into two sequences S₁ and S₂
 of n/2 elements each
 - Recur: recursively merge sort S₁ and S₂
 - Conquer: merge S₁ and S₂ into a sorted sequence

Merge Sort: Example

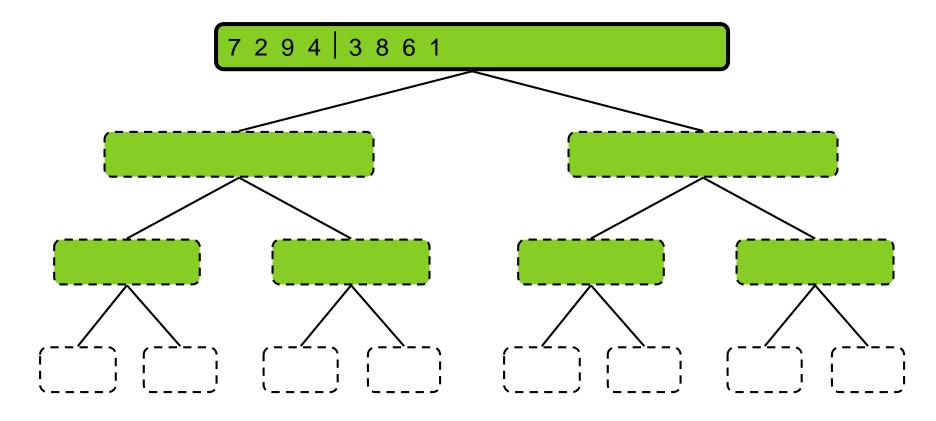
- The execution of merge sort can be depicted by a binary tree
 - each node represents a subproblem in merge sort, showing the unsorted subproblem before calling merge sort recursively
 - the root is the initial call
 - the leaves are the base cases of the recursion, with subsequences of size 0 or 1

Merge Sort: Example

Suppose we want to sort [7, 2, 9, 4, 3, 8, 6, 1]

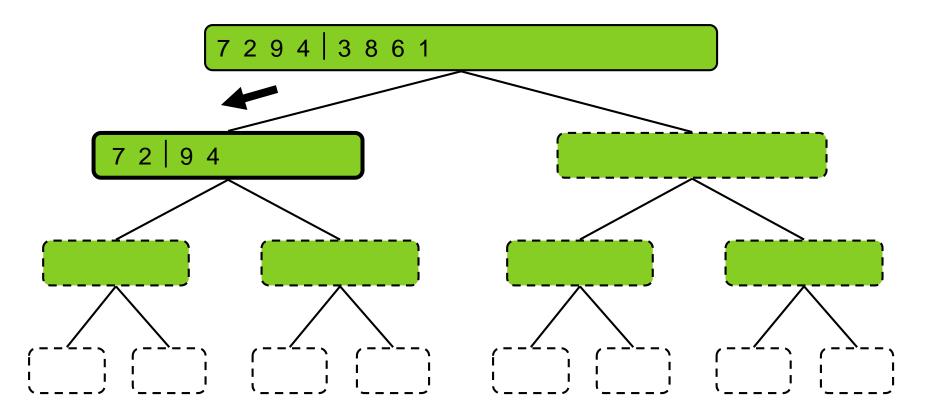
Merge Sort: Example (2)

Partition



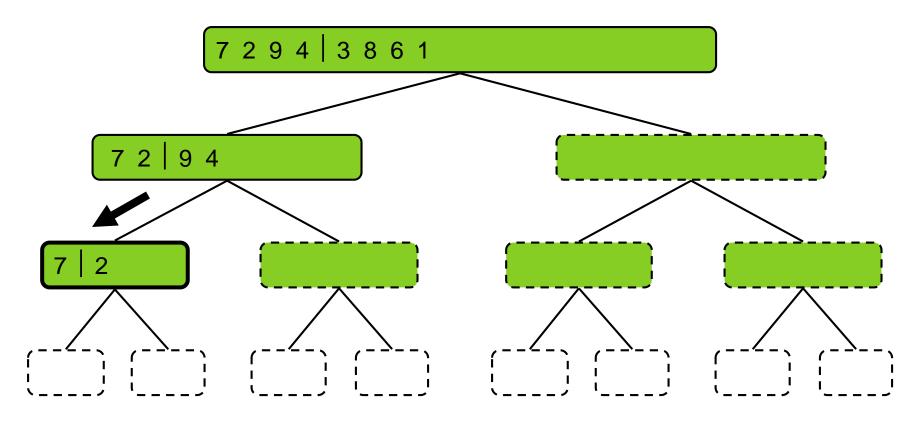
Merge Sort: Example (3)

Recursive call, partition



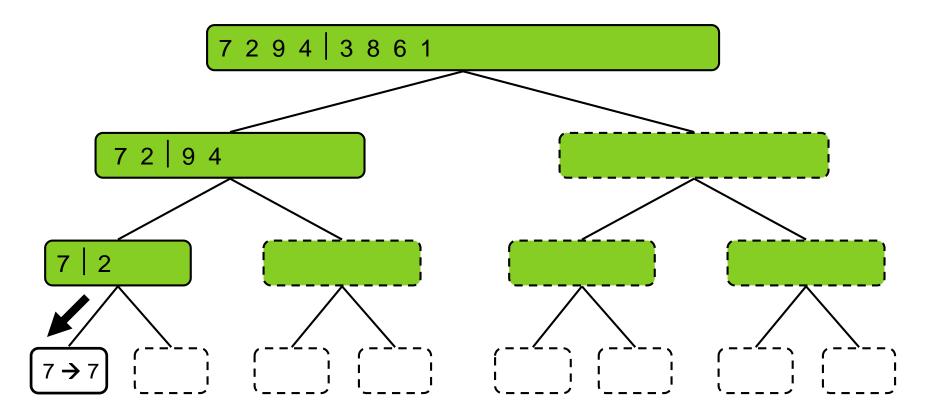
Merge Sort: Example (4)

Recursive call, partition



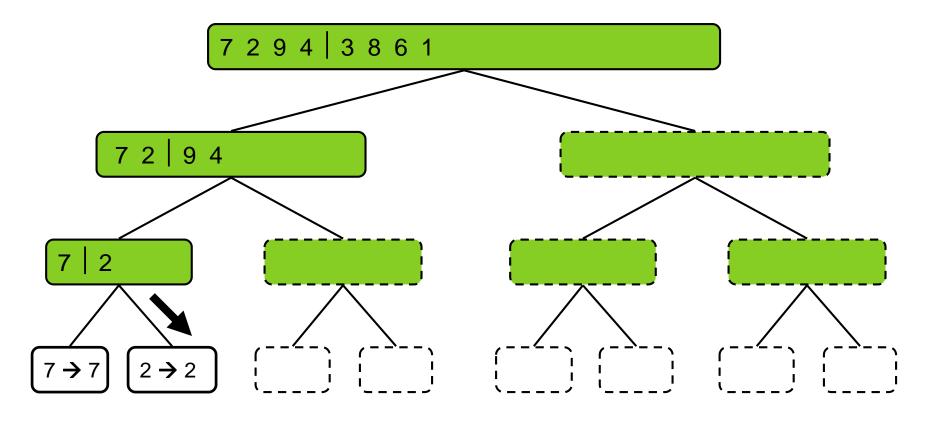
Merge Sort: Example (5)

Recursive call, base case



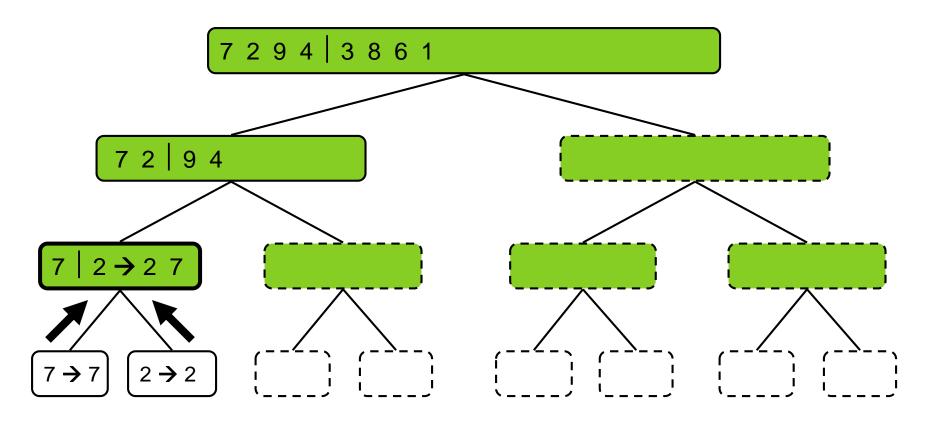
Merge Sort: Example (6)

Recursive call, base case



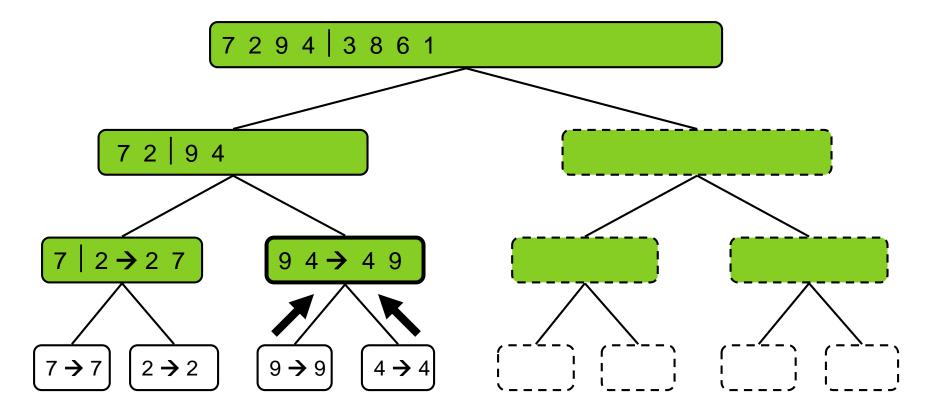
Merge Sort: Example (7)

Merge



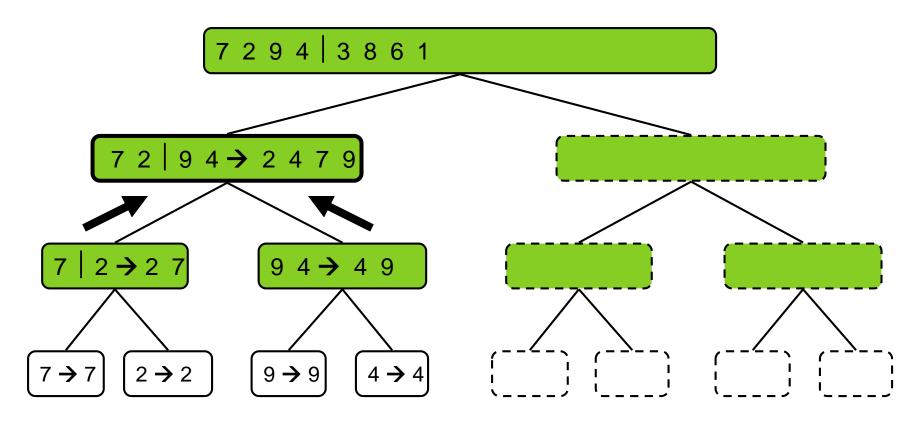
Merge Sort: Example (8)

Recursive call, ..., base case, merge



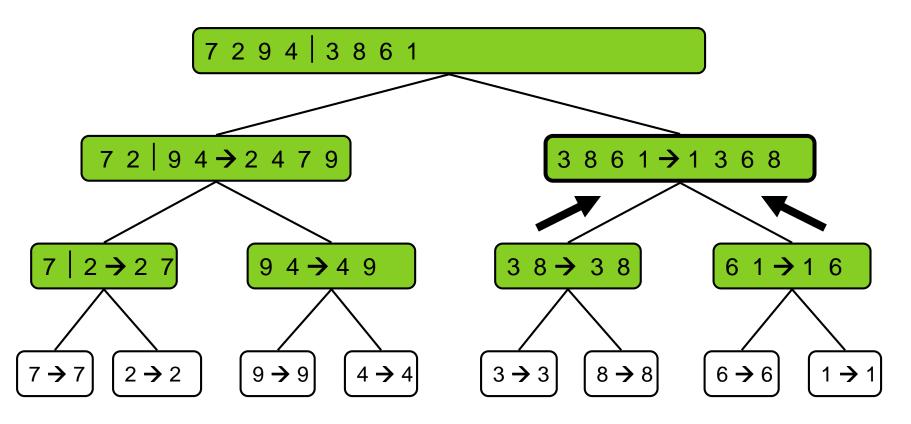
Merge Sort: Example (9)

Merge



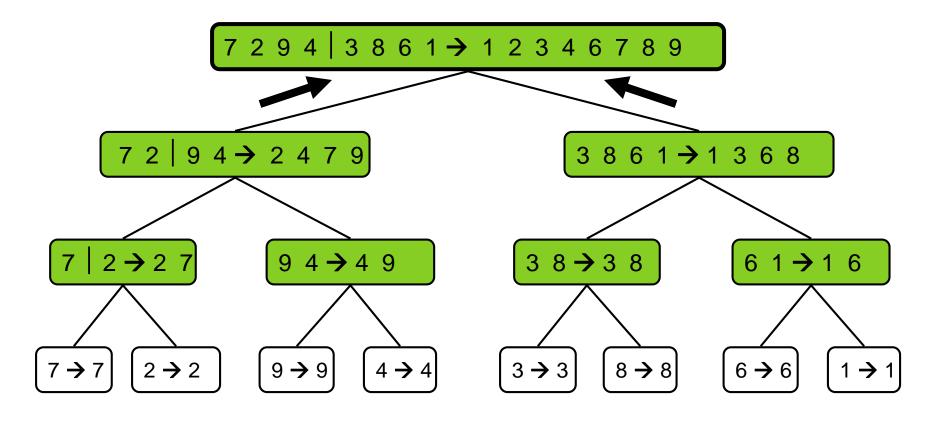
Merge Sort: Example (10)

Recursive call, ..., merge, merge



Merge Sort: Example (11)

Final merge and return



Merge Sort Pseudocode

```
function mergeSort(A):
   // Input: an unsorted array a
   // Output: array a in sorted order
   n = A.length
   if n <= 1:
      return A
   mid = n/2
   left = mergeSort(A[0 ... mid-1])
   right = mergeSort(A[mid ... n-1])
   return merge(left, right)
```

Merge Sort Pseudocode (2)

```
function merge(A, B):
   result = []
   aIndex = 0
   bIndex = 0
   while aIndex < A.length and bIndex < B.length:
      if A[aIndex] <= B[bIndex]:</pre>
         result.append(A[aIndex])
         aIndex++
      else:
          result.append(B[bIndex])
         bIndex++
   if aIndex < A.length:</pre>
      result = result + A[aIndex:end]
   if bIndex < B.length:</pre>
      result = result + B[bIndex:end]
   return result
```

Merge Sort Recurrence Relation

- Steps to merge sort:
 - Recursively merge sort the left half of the list
 - Recursively merge sort the right half of the list
 - 3. Merge both halves together
- Let T(n) be the running time of merge sort on an input of size n
- \cdot T(n) = step 1 + step 2 + step 3
- Notice that steps 1 and 2 are simply merge sorts on half the input and step 3 runs in O(n) time
- T(n) = T(n/2) + T(n/2) + O(n)= 2T(n/2) + O(n)

Merge Sort Recurrence Solution

- Recurrence relation:
 - Base case: $T(1) = c_1$
 - General case: T(n) = 2T(n/2) + O(n)
- Plug 'n' chug for a solution:

$$T(1) = c_1$$
 $= c_1$
 $T(2) = 2 T(1) + 2$ $= 2c_1 + 2$
 $T(4) = 2 T(2) + 4$ $= 2(2c_1 + 2) + 4$ $= 4c_1 + 8$
 $T(8) = 2 T(4) + 8$ $= 2(4c_1 + 8) + 8$ $= 8c_1 + 24$
 $T(16) = 2 T(8) + 16$ $= 2(8c_1 + 24) + 16$ $= 16c_1 + 64$
 $T(n)$ $= nc_1 + n \log n$
 $= O(n \log n)$

Analysis of Merge Sort

- To understand why merge sort is O(n log n), notice that the height h of the merge sort tree is O(log n) since it forms a perfect binary tree
- Overall amount of work done at each depth k is O(n) to partition and merge 2^k sequences of size n/2^k

| depth | sequences | size |
|-------|-----------|------------------|
| 0 | 1 | n |
| 1 | 2 | n/2 |
| 2 | 4 | n/4 |
| • | : | : |
| k | 2^k | n/2 ^k |

Solving Recurrence Relations

- To determine that merge sort was O(n log n), we determined the recurrence relation and used plug 'n' chug to conjecture a solution
 - $T(n) = T(n/2) + O(n) \rightarrow O(n \log n)$

Solving Recurrence Relations (2)

 Plug 'n' chug on recurrence relations is sort of a pain, but it turns out there's an easier way of solving recurrence relations...



 We will cover how to use Master Theorem, but the proof for why it works is in Dasgupta on pages 58-60

The Master Theorem

Where...

- a is the number of subproblems
- n/b is the size of each subproblem (if n/b is a fraction, b will be a whole number)
- n^d is the work done to prepare the subproblems and assemble the sub-results

Let $a \ge 1$, b > 1, $d \ge 0$, and T(n) be a monotonically increasing function of the form:

$$T(n) = aT(n/b) + \Theta(n^d)$$

Then:

$$\begin{array}{ll} T(n) \text{ is } \Theta(n^d) & \text{if } a < b^d \\ T(n) \text{ is } \Theta(n^d \log n) & \text{if } a = b^d \\ T(n) \text{ is } \Theta(n^{\log_b a}) & \text{if } a > b^d \end{array}$$

Applying the Master Theorem

$$T(n) = aT(n/b) + \Theta(n^d)$$

$$T(n) \text{ is } \Theta(n^d) \qquad \text{if } a < b^d$$

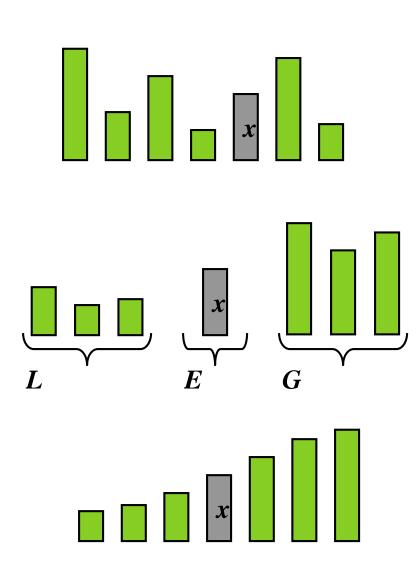
$$T(n) \text{ is } \Theta(n^d \log n) \qquad \text{if } a = b^d$$

$$T(n) \text{ is } \Theta(n^{\log_b a}) \qquad \text{if } a > b^d$$

- Merge sort's recurrence relation is
 T(n) = 2T(n/2) + O(n¹)
 - So, a = 2, b = 2, d = 1 and therefore $a = b^d$
 - T(n) is $\Theta(n^d \log n) = \Theta(n^1 \log n) = \Theta(n \log n)$

Quick Sort

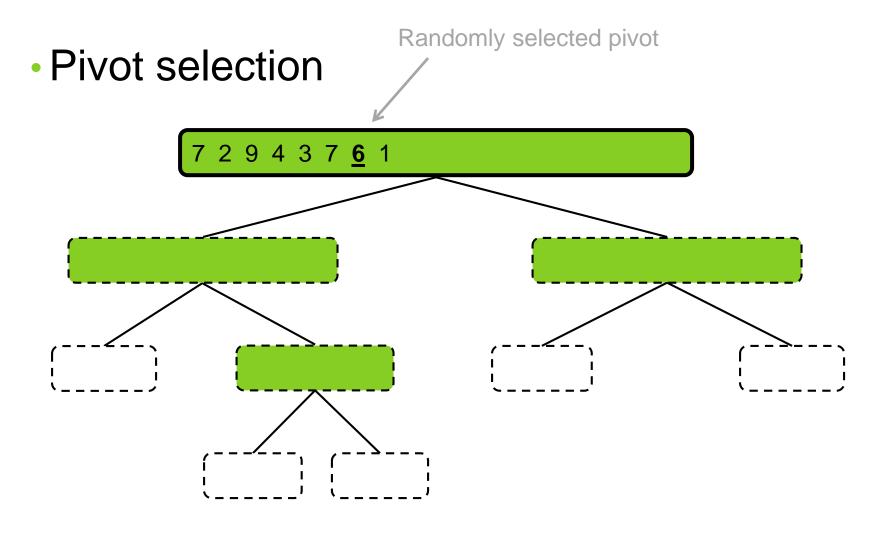
- Quick sort is a randomized sorting algorithm based on the divide-and-conquer paradigm:
 - Divide: pick a random element x (called the pivot) and partition sequence S into
 - L elements less than x
 - E elements equal to x
 - G elements greater than x
 - · Recur: quicksort L and G
 - Conquer: join L, E and G



Quick Sort: Example

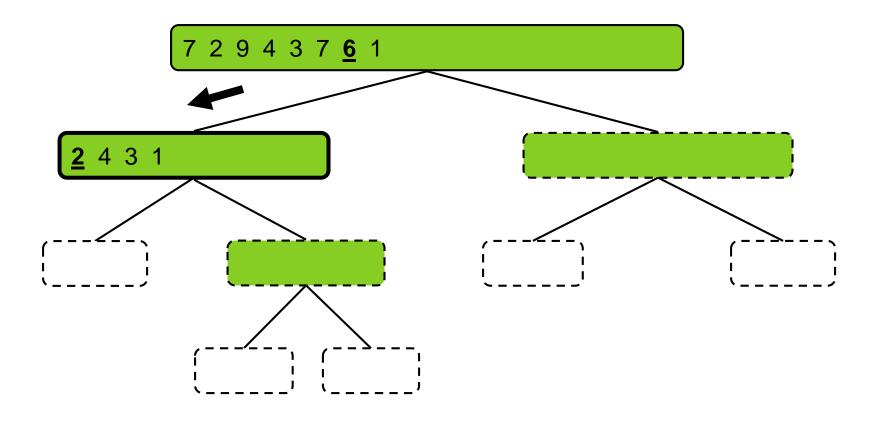
Suppose we want to sort [7, 2, 9, 4, 3, 8, 6, 1]

Quick Sort: Example



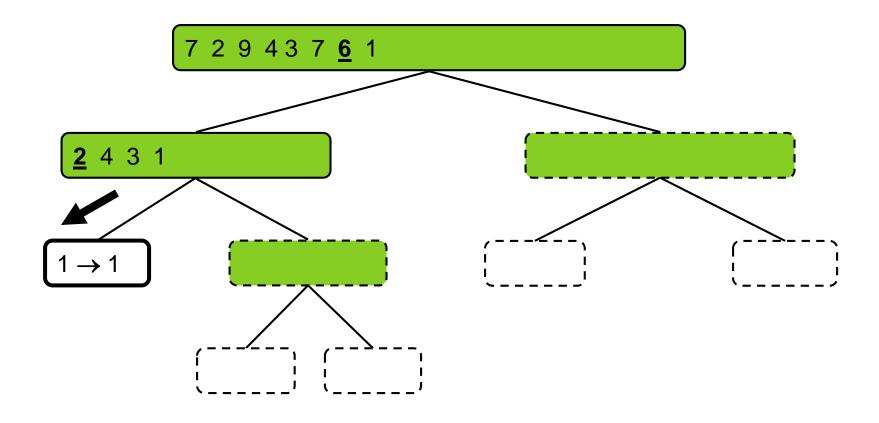
Quick Sort: Example (2)

Partition, recursive call, pivot selection



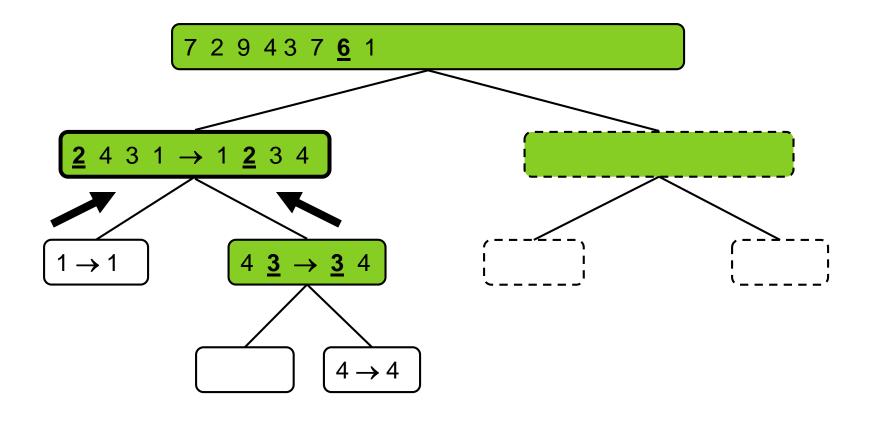
Quick Sort: Example (3)

Partition, recursive call, base case



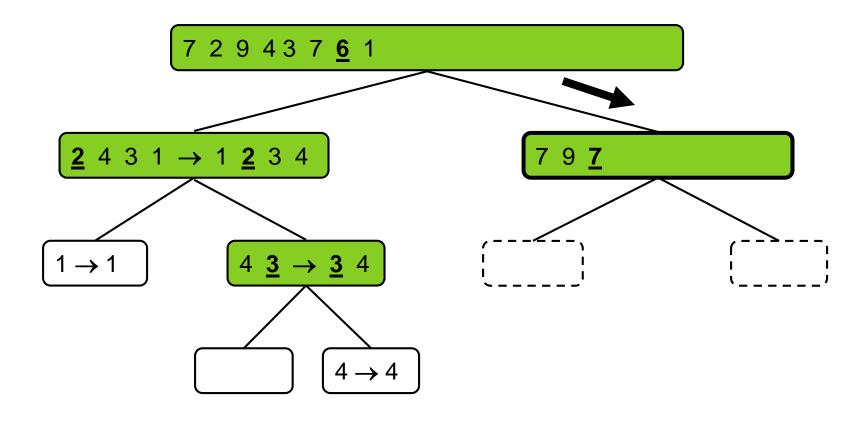
Quick Sort: Example (4)

Recursive call, ..., base case, join



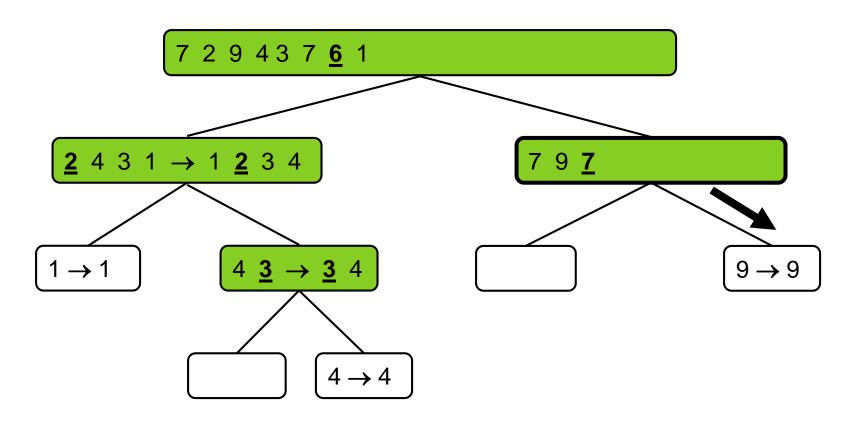
Quick Sort: Example (5)

Recursive call, pivot selection



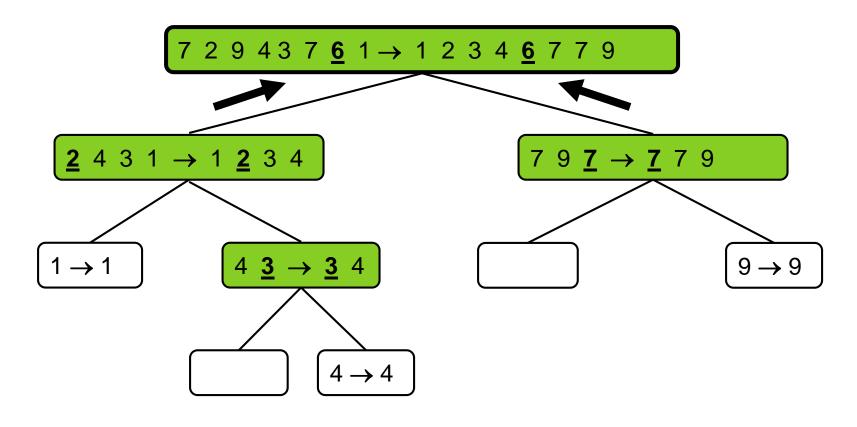
Quick Sort: Example (6)

Partition, ..., recursive call, base case



Quick Sort: Example (7)

Join, join



Quick Sort Pseudocode

```
function quick_sort(A):
   // Input: unsorted list
   // Output: sorted list
   if A.length ≤ 1
      return A
   pivot = random element from A
   L = [], E = [], G = []
   for each x in A:
      if x < pivot:</pre>
         L.append(x)
      else if x > pivot:
         G.append(x)
      else E.append(x)
   return quick_sort(L) + E + quick_sort(G)
```

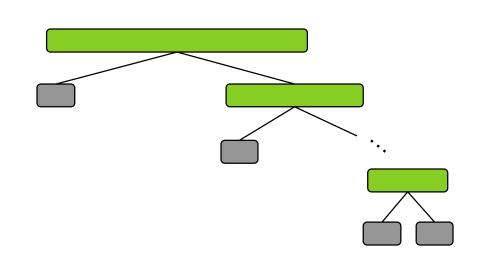
Worst-case Running Time

- The worst case for quick-sort occurs when the pivot is a unique minimum or maximum element
- One of L and G has size n 1 and the other has size 0
- The running time is proportional to the sum

$$n + (n - 1) + ... + 2 + 1$$

• Thus, the worst-case running time of quick-sort is $O(n^2)$

| depth | time |
|-------|-------|
| 0 | n |
| 1 | n - 1 |
| 2 | n-2 |
| : | : |
| n - 1 | 1 |



Expected-case Running Time

- To find an upper bound on the expected running time, assume there are no duplicates (if there are duplicates, the recursive calls will be even smaller)
- There are n possible unique ways quick sort will make its recursive calls:
 - |L| = 0, |G| = n-1
 - |L| = 1, |G| = n-2 :
 - |L| = n-1, |G| = 0
- Since there are n possible ways quick sort will recur, each has probability 1/n.
- We average over all possible splits and note that an additional linear amount of work is done:

$$T(n) = n + \frac{1}{n} \sum_{i=0}^{n-1} \left(T(i) + T(n-1-i) \right)$$

• The solution to this recurrence relation is (just trust us):

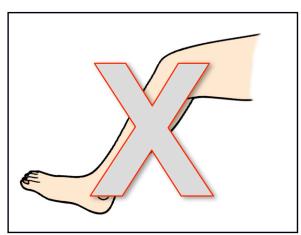
$$T(n) = 2n \ln n = 1.39n \log_2 n = O(n \log n)$$

In-Place Quick Sort?

```
function quick_sort(A):
   // Input: unsorted list
   // Output: sorted list
   if A.length ≤ 1
      return A
   pivot = random element from A
   L = [], E = [], G = []
   for each x in A:
      if x < pivot:</pre>
         L.append(x)
      else if x > pivot:
```

G.append(x)

else E.append(x)



Hmmm... This whole "LEG" business doesn't seem very in-place!

Let's see if we can partition the array in-place!

```
return quick sort(L) + E + quick sort(G)
```

In-Place Quick Sort!

```
function partition(A, low, high):
   // Input: list A with bounds low and high for the area
             we're partitioning
   // Output: The index of the pivot, once the array has been
              partitioned
   pivotIndex = random index between low and high
   pivotValue = A[pivotIndex]
   swap A[pivotIndex] and A[high] // move the pivot to end
   currIndex = low
   for i from low to high - 1:
       if A[i] <= pivotValue :</pre>
           swap A[i] and A[currIndex]
           currIndex++
   swap A[currIndex] and A[high] // move the pivot back
   return currIndex
```

In-Place Quick Sort!

```
function quick_sort(A, low, high):
    // Input: unsorted list
    // Output: sorted list between low and high
    if low < high:
        pivotIndex = partition(A, low, high)
        quicksort(A, low, pivotIndex - 1)
        quicksort(A, pivotIndex + 1, high)</pre>
```

How fast can we sort?

- Both merge and quick sort are O(n log n)
- Can we do better? No! (sort of)

- Claim: Any comparison sort must in the worst case make at least $\Omega(n \log n)$ comparisons to sort n keys.
- Let's try to prove this!

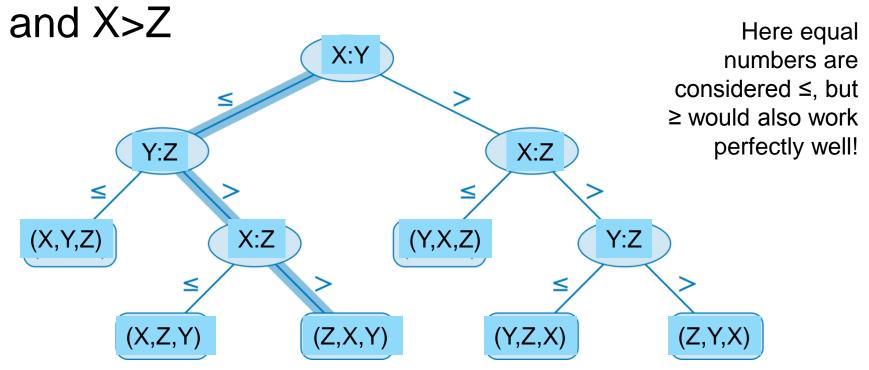
Comparative Sorting is Ω(n log n)

- A sorting algorithm, viewed abstractly, takes a sequence of keys k₁, k₂, ... k_n and outputs the permutation that will sort them.
- For a given n, we can represent the optimal algorithm as a perfect binary decision tree:
 - internal nodes are comparisons of two keys
 - leaves are the correct permutations.
- Sorting a particular sequence is equivalent to walking the tree from the root to a leaf. The number of comparisons in the worst case is the height of the tree.

Comparison Tree

 The path to any leaf shows the minimum number of comparisons necessary

If (Z,X,Y) is the proper sort, then X≤Y, Y>Z,



Comparative Sorting Proof (2)

- n! permutations of a given sequence, so the tree has n! leaves.
- We know that a perfect binary tree with height h has 2^h leaves. Thus, a tree with n! leaves has height log (n!)
- Stirling's Formula:

```
n! \geq n^n e^{-n}

log(n!) \geq log(n^n e^{-n})

log(n!) \geq n log n - n log e
```

• Therefore the height (and number of comparisons in the worst case) is $\Omega(n \log n)$.

Non-comparative Sorting

- Sorting functions may need to accept a variety of inputs
 - Integers
 - Floats
 - Strings
 - Arrays
 - Other objects... People? Cows?
- As long as there is some way to compare elements in the input domain, we can apply one of the comparison-based sorting algorithms
- But sometimes, if we know something about our input domain in advance, we be a little cleverer in our sorting...
- Radix sort: great for positive integers!

Non-comparative Sorting (2)

- First let's think about an easier problem:
 - Given an array of integers between 0 and 9, sort the array
 - e.g. [5, 1, 6, 2, 3, 1] -> [1, 1, 2, 3, 5, 6]
- Since we know that every element is guaranteed to be an integer between 0 and 9, let's use that to our advantage!
 - Make an array of 10 buckets, one for each possible number
 - For each number x in the array, add it to the bucket at index x
 - Return the concatenation of all the buckets, in order

$$[1, 1] + [2] + [3] + [5] + [6]$$

What's the runtime of this? O(n)!

Radix Sort

- What about sorting integers of unknown size?
- Check out these two numbers258391 258492
- How would you compare these by hand?
 - Digit by digit, probably!
 - The 3 highest order bits are the same (258), so we keep going until we find that 4 is greater than 3, so 258492 must be greater than 258391
- Radix sort takes advantage of the digity-ness of integers, and also the fact that for any given digit, there are a constant number of options (0-9)

- Goal: Sort the list [273, 279, 8271, 7891, 8736, 8735]
- Start with the lowest order digit (the 1's place) and add each number to the bucket corresponding to that digit

| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
|---|------|---|-----|---|------|------|---|---|-----|
| | 8271 | | 273 | | 8735 | 8736 | | | 279 |
| | 7891 | | | | | | | | |

- After sorting, concatenate all buckets in order to get a new list: [8271, 7891, 273, 8735, 8736, 279]
- Everything's now sorted by 1's place

- The list is now [8271, 7891, 273, 8735, 8736, 279]
- Now we repeat, sorting by 10's place

| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
|---|---|---|------|---|---|---|------|---|------|
| | | | 8735 | | | | 8271 | | 7891 |
| | | | 8736 | | | | 273 | | |
| | | | | | | | 279 | | |

- New list: [8735, 8736, 8271, 273, 279, 7891]
- Everything's now sorted by 10's and 1's places

- The list is now [8735, 8736, 8271, 273, 279, 7891]
- Now we repeat, sorting by 100's place

| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
|---|---|------|---|---|---|---|------|------|---|
| | | 8271 | | | | | 8735 | 7891 | |
| | | 273 | | | | | 8736 | | |
| | | 279 | | | | | | | |

- New list: [8271, 273, 279, 8735, 8736, 7891]
- Everything's now sorted by 100's, 10's and 1's places

- The list is now [8271, 273, 279, 8735, 8736, 7891]
- Now we repeat, sorting by 1000's place
 - We treat numbers with fewer than 4 digits as if they had additional leading 0's

| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
|-----|---|---|---|---|---|---|------|------|---|
| 273 | | | | | | | 7891 | 8271 | |
| 279 | | | | | | | | 8735 | |
| | | | | | | | | 8736 | |

- New list: [273, 279, 7891, 8271, 8735, 8736]
- We've finished sorting by all digits! The list is sorted!

Radix Sort

```
function radix sort(A):
   //Input: unsorted list of positive integers
  //Output: sorted list
   buckets = array of 10 lists
   for place = least to most significant
      for number in A
         d = digit in number at place
         buckets[d].append(number)
      A = concatenate all buckets in order
      empty all buckets
   return A
```

 Very efficient: O(n*d), where d is the number of digits in the largest number.

Radix Sort

- Radix sort can also be applied to different kinds of inputs, other than positive integers in base-10
 - Octal (base-8) or hexadecimal (base-16) numbers
 - Strings: make 1 bucket for each valid letter or character
- The number of buckets don't need to be the same for each "round" of sorting
 - Sorting a deck of cards: use numbers as the first buckets (lowest order bit) and suits as second buckets (highest order bit)
- You can also represent just about anything as a binary sequence of 0's and 1's and radix sort using two buckets
 - But then the number of digits will dominate the runtime, and for very long sequences, that sucks

Summary of Sorting Algorithms

| Algorithm | Time | Notes |
|----------------|------------------------|--------------------------------------------------------------------------------|
| Selection sort | O(n ²) | in-place slow (good for small inputs) |
| Insertion sort | O(n ²) | in-place slow (good for small inputs) |
| Merge sort | O(n log n) | fast (good for large inputs) |
| Quick sort | O(n log n) expected | randomized fastest (good for large inputs) |
| Radix sort | O(nd) | d is number of digits in largest number basically linear when d is small |

Readings

- Dasgupta
 - Section 2.1: A good complementary introduction to divide and conquer algorithms
 - Section 2.2: A review of recurrence relations and the master theorem
 - Section 2.3: Analysis of merge sort and a lower bound on comparative sorting

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