Data Structures and Algorithms



Outline

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- Merge Sort
- Quick Sort
- Sorting Lower Bound
- Bucket-Sort
- Radix Sort

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Merge Sort



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 S₁ and S₂
 - Conquer: combine the solutions for S₁ and S₂ into a solution for S
- The base case for the recursion are subproblems of size 0 or 1
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- Merge-sort is a sorting algorithm based on the divide-and-conquer paradigm
- Like heap-sort
 - It uses a comparator
 - It has O(n log n) running time
- Unlike heap-sort
 - It does not use an auxiliary priority queue
 - It 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 S₁ and S₂ of about n/2 elements each
 - Recur: recursively sort S₁ and S₂
 - Conquer: merge S₁ and S₂ 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)

 $S \leftarrow merge(S_1, S_2)$

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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 elements and implemented by means of a doubly linked list, takes O(n) time

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Algorithm merge(A, B)
Input sequences A and B with n/2 elements each
Output sorted sequence of A ∪ B

 $S \leftarrow$ empty sequence

while ¬A.isEmpty() ∧ ¬B.isEmpty()
if A.first().element() < B.first().element()
S.insertLast(A.remove(A.first()))</pre>

else

S.insertLast(B.remove(B.first())) while ¬A.isEmpty() S.insertLast(A.remove(A.first())) while ¬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 0 or 1























Analysis of Merge-Sort

- The height h of the merge-sort tree is $O(\log n)$
 - at each recursive call we divide in half the sequence,
- 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 $O(n \log n)$



Summary of Sorting Algorithms

Algorithm	Time	Notes	
selection-sort	O (n ²)	 ◆ slow ◆ in-place ◆ for small data sets (< 1K) 	
insertion-sort	O (n ²)	 slow in-place for small data sets (< 1K) 	
heap-sort	O (n log n)	 ◆ fast ◆ in-place ◆ for large data sets (1K — 1M) 	
merge-sort	O (n log n)	 fast sequential data access for huge data sets (> 1M) 	
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Nonrecursive Merge-Sort

- public static void mergeSort(Object[] orig, Comparator c) { //
 nonrecursive
 Object[] in = new Object[orig.length]; // make a new temporary array
 System.arraycopy(orig,0,in,0,in.length); // copy the input
 Object[] out = new Object[in.length]; // output array
 Object[] temp; // temp array reference used for swapping
 int n = in.length;
 - for (int i=1; i < n; i*=2) { // each iteration sorts all length-2*i runs
 for (int j=0; j < n; j+=2*i) // each iteration merges two length-i pairs
 merge(in,out,c,j,i); // merge from in to out two length-i runs at j
 temp = in; in = out; out = temp; // swap arrays for next iteration</pre>

// the "in" array contains the sorted array, so re-copy it System.arraycopy(in,0,orig,0,in.length);

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Nonrecursive Merge-Sort

else if (y < end2) // second run didn't finish System.arraycopy(in, y, out, z, end2 - y);

merge runs of length 2, then 4, then 8, and so on

merge two runs in the in array to the out array

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public static void mergeSort(Object[] orig, Comparator c) { // nonrecursive Object[] in = new Object[orig.length]; // make a new temporary array System.arraycopy(orig,0,in,0,in.length); // copy the input Object[] out = new Object[in.length]; // output array Object[] temp; // temp array reference used for swapping int n = in.length;for (int i=1; i < n; i*=2) { // each iteration sorts all length-2*i runs for (int j=0; j < n; j+= 2^{*i}) // each iteration merges two length-i pairs merge(in,out,c,j,i); // merge from in to out two length-i runs at j temp = in; in = out; out = temp; // swap arrays for next iteration // the "in" array contains the sorted array, so re-copy it System.arraycopy(in,0,orig,0,in.length); protected static void merge(Object[] in, Object[] out, Comparator c, int start, int inc) { // merge in[start..start+inc-1] and in[start+inc..start+2*inc-1] int x = start; // index into run #1 int end1 = Math.min(start+inc, in.length); // boundary for run #1 int end2 = Math.min(start+2*inc, in.length); // boundary for run #2 int y = start+inc; // index into run #2 (could be beyond array boundary) int z =start; // index into the out array while ((x < end1) && (y < end2))if $(c.compare(in[x],in[y]) \le 0)$ out[z++] = in[x++];else out[z++] = in[y++];if (x < end1) // first run didn't finish System.arraycopy(in, x, out, z, end1 - x);

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Quick-Sort



Quick-Sort

L

X

G

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E

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



Partition

- We partition an input sequence as follows:
 - We remove, in turn, each element y from S and
 - We insert y into L, E or G, depending on the result of the comparison with the pivot x
- Each insertion and removal is at the beginning or at the end of a sequence, and hence takes **O**(1) time
- Thus, the partition step of quick-sort takes O(n) time

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Algorithm *partition*(*S*, *p*) Input sequence S, position p of pivot Output subsequences L, E, G of the elements of S less than, equal to, or greater than the pivot, resp. *L*, *E*, *G* \leftarrow empty sequences $x \leftarrow S.remove(p)$ while ¬*S.isEmpty*() $y \leftarrow S.remove(S.first())$ if y < x*L.insertLast(y)* else if y = x*E.insertLast(y)* else $\{y > x\}$ *G.insertLast(y)* return L, E, G

Quick-Sort Tree

- An execution of quick-sort is depicted by a binary tree
 - Each node represents a recursive call of quick-sort and stores
 - Unsorted sequence before the execution and its pivot
 - Sorted sequence at the end of the execution
 - The root is the initial call
 - The leaves are calls on subsequences of size 0 or 1







· Partition, recursive call, pivot selection













Worst-case Running Time

- The worst case for quick-sort occurs when the pivot is the 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

 $n + (n - 1) + \dots + 2 + 1$

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

depth time



Expected Running Time

- Consider a recursive call of quick-sort on a sequence of size s
 - Good call: the sizes of L and G are each less than 3s/4
 - Bad call: one of L and G has size greater than 3s/4



Expected Running Time, Part 2

- Probabilistic Fact: The expected number of coin tosses required in order to get k heads is 2k
- For a node of with input size s, the input sizes of its children are each at most s3/4 or s/(4/3).



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In-Place Quick-Sor

- Quick-sort can be implemented to run in-place
- In the partition step, we use replace operations to rearrange the elements of the input sequence such that
 - the elements less than the pivot have rank less than h
 - the elements equal to the pivot have rank between h and k
 - the elements greater than the pivot have rank greater than k
- The recursive calls consider
 - elements with rank less than h
 - elements with rank greater than k

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Algorithm *inPlaceQuickSort*(*S*, *l*, *r*)

Input sequence *S*, ranks *l* and *r* Output sequence *S* with the

elements of rank between *l* and *r* rearranged in increasing order

if $l \ge r$

return

- $i \leftarrow$ a random integer between l and r
- $x \leftarrow S.elemAtRank(i)$
- $(h, k) \leftarrow inPlacePartition(x)$

inPlaceQuickSort(S, l, h - 1)

inPlaceQuickSort(S, k + 1, r)

In-Place Partitioning

 Perform the partition using two indices to split S into L and E U G (a similar method can split E U G into E and G).
 i

3 2 5 1 0 7 3 5 9 2 7 9 8 9 7 <u>6</u> 9

(pivot = 6)

- Repeat until j and k cross:
 - Scan j to the right until finding an element $\geq x$.
 - Scan k to the left until finding an element < x.
 - Swap elements at indices j and k

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)
quick-sort	O(n log n) expected	 in-place, randomized fastest (good for large inputs)
heap-sort	O (n log n)	 in-place fast (good for large inputs)
merge-sort	O (n log n)	 sequential data access fast (good for huge inputs)
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Java Implementation

public static void quickSort (Object[] S, Comparator c) {
 if (S.length < 2) return; // the array is already sorted in this case
 quickSortStep(S, c, 0, S.length-1); // recursive sort method</pre>

only works for distinct elements private static void quickSortStep (Object[] S, Comparator c, int leftBound, int rightBound) { if (leftBound >= rightBound) return; // the indices have crossed Object temp; // temp object used for swapping Object pivot = S[rightBound]; int leftIndex = leftBound; // will scan rightward int rightIndex = rightBound-1; // will scan leftward while (leftIndex <= rightIndex) { // scan right until larger than the pivot while ((leftIndex <= rightIndex) && (c.compare(S[leftIndex], pivot)<=0)) leftIndex++: // scan leftward to find an element smaller than the pivot while ((rightIndex \geq leftIndex) && (c.compare(S[rightIndex], pivot)>=0)) rightIndex--; if (leftIndex < rightIndex) { // both elements were found temp = S[rightIndex];S[rightIndex] = S[leftIndex]; // swap these elements S[leftIndex] = temp; } // the loop continues until the indices cross temp = S[rightBound]; // swap pivot with the element at leftIndex S[rightBound] = S[leftIndex];S[leftIndex] = temp; // the pivot is now at leftIndex, so recurse quickSortStep(S, c, leftBound, leftIndex-1); quickSortStep(S, c, leftIndex+1, rightBound);

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Sorting Lower Bound



Comparison-Based Sorting



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- Many sorting algorithms are comparison based.
 - They sort by making comparisons between pairs of objects
 - Examples: bubble-sort, selection-sort, insertion-sort, heap-sort, merge-sort, quick-sort, ...
- Let us therefore derive a lower bound on the running time of any algorithm that uses comparisons to sort n elements, x₁, x₂, ..., x_n.



Counting Comparisons

 $x_c < x_d$?

 $\left(x_{m} < x_{o}?\right) \left(x_{p} < x_{q}?\right)$

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· Let us just count comparisons then.

 $x_a < x_b$?

 $\left[x_e < x_f?\right] \left[x_k < x_l?\right]$

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• Each possible run of the algorithm corresponds to a root-to-leaf path in a decision tree $x_i < x_j$?

Decision Tree Height

- The height of this decision tree is a lower bound on the running time
- Every possible input permutation must lead to a separate leaf output.
 - If not, some input ...4...5... would have same output ordering as ...
 5...4..., which would be wrong.
- Since there are n!=1*2*...*n leaves, the height is at least log (n!)



The Lower Bound



Therefore, any such algorithm takes time at least

$$\log(n!) \ge \log\left(\frac{n}{2}\right)^{\frac{n}{2}} = (n/2)\log(n/2)$$

 That is, any comparison-based sorting algorithm must run in Ω(n log n) time.

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Bucket-Sort and Radix-Sort



Bucket-Sort



- Bucket-sort uses the keys as indices into an auxiliary array *B* of sequences (buckets)
 - Phase 1: Empty sequence *S* by moving each entry (*k*, *o*) into its bucket *B*[*k*]
 - Phase 2: For i = 0, ..., N-1, move the entries of bucket B[i] to the end of sequence S

Analysis:

- Phase 1 takes O(n) time
- Phase 2 takes O(n + N) time

Bucket-sort takes O(n + N) time Phạm Bảo Sơn DSA Algorithm *bucketSort*(S, N) Input sequence S of (key, element) items with keys in the range [0, N-1]**Output** sequence *S* sorted by increasing keys $B \leftarrow$ array of N empty sequences while ¬*S.isEmpty(*) $f \leftarrow S.first()$ $(k, o) \leftarrow S.remove(f)$ B[k].insertLast((k, o)) for $i \leftarrow 0$ to N - 1while ¬*B*[*i*].*isEmpty*() $f \leftarrow B[i].first()$ $(k, o) \leftarrow B[i]$.remove(f) S.insertLast((k, o))



Properties and Extension

- Key-type Property
 - The keys are used as indices into an array and cannot be arbitrary objects
 - No external comparator
- Stable Sort Property
 - The relative order of any two items with the same key is preserved after the execution of the algorithm

Extensions

- Integer keys in the range [a, b]
 - Put entry (k, o) into bucket
 B[k a]
- String keys from a set *D* of possible strings, where *D* has constant size (e.g., names of the 50 U.S. states)
 - Sort *D* and compute the rank
 r(*k*) of each string *k* of *D* in the sorted sequence
 - Put entry (k, o) into bucket
 B[r(k)]

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Lexicographic Orde

- A *d*-tuple is a sequence of *d* keys (*k*₁, *k*₂, ..., *k_d*), where key *k_i* is said to be the *i*-th dimension of the tuple
- Example:
 - The Cartesian coordinates of a point in space are a 3-tuple
- The lexicographic order of two *d*-tuples is recursively defined as follows

 $(x_1, x_2, ..., x_d) < (y_1, y_2, ..., y_d)$

 $x_1 < y_1 \lor x_1 = y_1 \land (x_2, ..., x_d) < (y_2, ..., y_d)$

I.e., the tuples are compared by the first dimension, then by the second dimension, etc.

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Lexicographic-Sort

- Let *C_i* be the comparator that compares two tuples by their *i*-th dimension
- Let *stableSort*(*S*, *C*) be a stable sorting algorithm that uses comparator *C*
- Lexicographic-sort sorts a sequence of *d*-tuples in lexicographic order by executing *d* times algorithm *stableSort*, one per dimension
- Lexicographic-sort runs in *O*(*dT*(*n*)) time, where *T*(*n*) is the running time of *stableSort* Pham Bảo Sơn DSA

Algorithm *lexicographicSort(S)* Input sequence S of d-tuples Output sequence S sorted in lexicographic order

for $i \leftarrow d$ downto 1 stableSort(S, C_i)

Example:

(7,4,6) (5,1,5) (2,4,6) (2, 1, 4) (3, 2, 4)

(2, 1, 4) (3, 2, 4) (5, 1, 5) (7, 4, 6) (2, 4, 6)

(2, 1, 4) (5, 1, 5) (3, 2, 4) (7, 4, 6) (2, 4, 6)

(2, 1, 4) (2,4,6) (3, 2, 4) (5,1,5) (7,4,6)

Radix-Sort

- Radix-sort is a specialization of lexicographic-sort that uses bucket-sort as the stable sorting algorithm in each dimension
- Radix-sort is applicable to tuples where the keys in each dimension *i* are integers in the range [0, *N*-1]
- Radix-sort runs in time O(d(n + N))

Algorithm radixSort(S, N) Input sequence S of d-tuples such that $(0, ..., 0) \le (x_1, ..., x_d)$ and $(x_1, ..., x_d) \le (N-1, ..., N-1)$ for each tuple $(x_1, ..., x_d)$ in S Output sequence S sorted in lexicographic order for $i \leftarrow d$ downto 1 bucketSort(S, N)

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Radix-Sort for Binary Numbers

Consider a sequence of *n b*-bit integers

 $\boldsymbol{x} = \boldsymbol{x}_{\boldsymbol{b}-1} \dots \boldsymbol{x}_1 \boldsymbol{x}_0$

- We represent each element as a *b*-tuple of integers in the range [0, 1] and apply radix-sort with *N* = 2
- This application of the radix-sort algorithm runs in O(bn) time
- For example, we can sort a sequence of 32-bit integers in linear time Pham Bảo Sơn DSA

Algorithm *binaryRadixSort(S)*

Input sequence S of b-bit integers Output sequence S sorted replace each element xof S with the item (0, x)for $i \leftarrow 0$ to b - 1

replace the key k of each item (k, x) of S with bit x_i of x bucketSort(S, 2)

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Example

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Sorting a sequence of 4-bit integers

