# Probabilistic Algorithms: What, Why, and How

A Deep Dive into Randomness in Computing

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## **Outline**

What are Probabilistic Algorithms?

Applications of Probabilistic Algorithms

Why Probabilistic Algorithms?

How do Probabilistic Algorithms Work?

Example: Randomized Quicksort

Step-by-Step Execution

Time Complexity Analysis of Randomized Quicksort

Example: Monte Carlo Estimation of

Probabilistic Data Structures

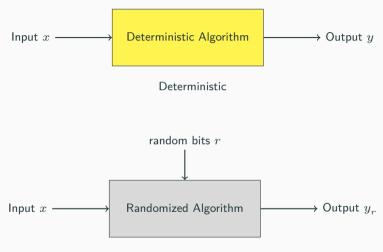
What are Probabilistic Algorithms?

## What is a Probabilistic Algorithm?

## **Definition**

An algorithm that makes random choices during execution to influence its behavior or output.

## **Deterministic vs Probabilistic Algorithm**



Probabilistic

# Types of Probabilistic Algorithms

- Las Vegas Algorithms (Babai, 1979)
- Monte Carlo Algorithms (Metropolis & and, 1949)

## Las Vegas Algorithms

**Definition:** A Las Vegas algorithm always produces a correct result or reports failure, with the running time depending on random choices. (Gupta & Ramachandran, 1992)

## **Monte Carlo Algorithms**

**Definition:** A Monte Carlo algorithm has a probability of producing an incorrect result, but its running time is bounded. (James, 1990)

**Applications of Probabilistic** 

**Algorithms** 

## **Real-World Motivation**

- Web search (PageRank)
- Load balancing (power of two choices)
- Hashing (universal hash functions)
- Primality testing (Miller-Rabin)

Why Probabilistic Algorithms?

## Why Randomness?

## Motivation

- Simpler algorithms
- Better expected performance
- Avoid worst-case scenarios
- Useful for large-scale and distributed systems

# How do Probabilistic Algorithms

Work?

## **How: Randomization in Algorithms**

## **Key Idea**

Use random choices to influence the algorithm's path or output.

- Random pivot in Quicksort
- Random walks in graphs
- Random sampling

**Example: Randomized Quicksort** 

## QuickSort vs Randomized QuickSort

## QuickSort:

- 1. Pick a pivot element from the array (Hoare, 1962)
- 2. Split array into 3 subarrays: those smaller than pivot, those larger than pivot, and the pivot itself
- 3. Recursively sort the subarrays, and concatenate them

## Randomized QuickSort:

- Pick a pivot element uniformly at random from the array (Motwani & Raghavan, 1995)
- 2. Split array into 3 subarrays: those smaller than pivot, those larger than pivot, and the pivot itself
- 3. Recursively sort the subarrays, and concatenate them

## **Example: Randomized Quicksort**

**Recall:** QuickSort can take  $\Omega(n^2)$  time to sort an array of size n (Sedgewick, 1978)

## Randomized QuickSort: Expected Runtime

## **Theorem**

Randomized QuickSort sorts a given array of length n in  $O(n\log n)$  expected time. (Sedgewick, 1977)

**Note:** On every input, randomized QuickSort takes  $O(n\log n)$  time in expectation. On every input, it may take  $\Omega(n^2)$  time with some small probability.

# Randomized Quicksort: Step 1 (Initial Array)

Consider the array:

15	3	1	10	9	0	6	4
----	---	---	----	---	---	---	---

## Randomized Quicksort: Step 1.1 (Pivot Chosen)

Suppose the random pivot chosen is 10 (at index 3):

15	3	1	10	9	0	6	4
----	---	---	----	---	---	---	---

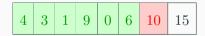


## Randomized Quicksort: Step 2 (Partitioning Around Pivot 10)

After selecting pivot 10, we partition the array:

- Left: 4, 3, 1, 9, 0, 6 (elements before pivot position)
- Middle: 10 (pivot)
- Right: 15 (element after pivot position)

## After partitioning:

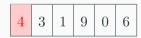


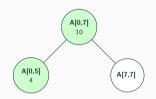


# Randomized Quicksort: Step 3 (Recurse Left [A[0,5]], Pivot 4)

Recurse on the left subarray:

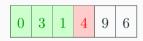
Let's choose a random pivot, say 4.





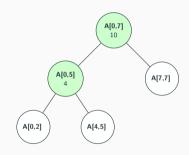
# Randomized Quicksort: Step 3.1 (Partition Left [A[0,5]] Around 4)

After partitioning the left subarray:



### Partition:

- Left: 0, 3, 1 (elements before pivot)
- Middle: 4 (pivot)
- Right: 9, 6 (elements after pivot)

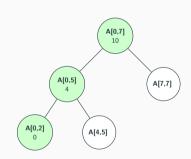


## Randomized Quicksort: Step 3.1.1 (Recurse Left [A[0,2]], Pivot 0)

Recurse on the left subarray:

Let's choose a random pivot, say 0.

0 3 1



# Randomized Quicksort: Step 3.1.1.1 (Partition Left [A[0,2]] Around 0)

After partitioning the left subarray:

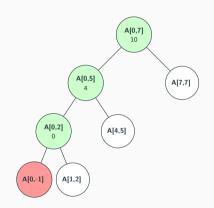


#### Partition:

Left: (empty)

Middle: 0 (pivot)

• Right: 3, 1 (elements after pivot)

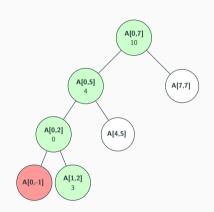


# Randomized Quicksort: Step 3.1.1.2 (Recurse Right [A[1,2]], Pivot 3)

Recurse on the right subarray:

Let's choose a random pivot, say 3.

3 1



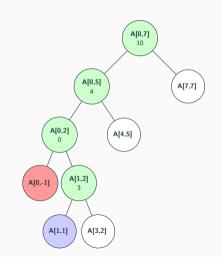
# Randomized Quicksort: Step 3.1.1.2.1 (Partition [A[1,2]] Around 3)

After partitioning the left subarray:



## Partition:

- Left: 1 (element before pivot)
- Middle: 3 (pivot)
- Right: (empty)



## Randomized Quicksort: Step 3.1.1.2.1.1 (Recurse Left [A[1,1]], Done)

After partitioning the left subarray:

1

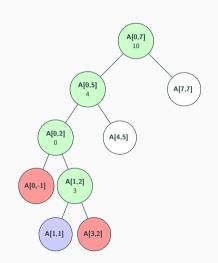
#### Partition:

• Left: (empty)

Middle: 1 (pivot)

Right: (empty)

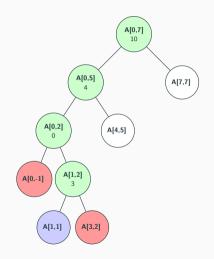
Single element subarray, done, return.



## Randomized Quicksort: Step 3.1.1.2.1.2 (Recurse Right [A[3,2]], Done)

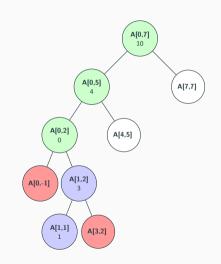
Recurse on the right subarray A[3,2] (empty, done).

Return to parent call A[1,2]



## Randomized Quicksort: Step 3.1.2 (Return to [A[0,2]])

Return to parent call A[0,2]

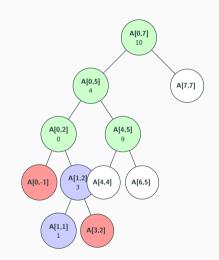


# Randomized Quicksort: Step 3.2 (Recurse Right [A[4,5]], Pivot 9)

Recurse on the right subarray A[4,5]:

Let's choose a random pivot, say 9.

9 6



# Randomized Quicksort: Step 3.2.1 (Partition [A[4,5]] Around 9)

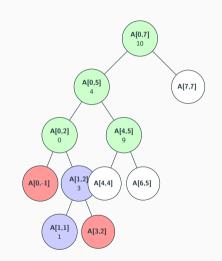
Recurse on the right subarray A[4,5]:

Suppose the random pivot is 9:

6 <mark>9</mark>

## Partition:

- Left: 6 (element before pivot)
- Middle: 9 (pivot)
- Right: (empty)



# Randomized Quicksort: Step 3.2.1.1 (Recurse Left [A[4,4]], Done)

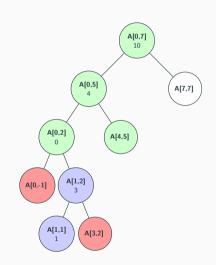
After partitioning the left subarray:

6

## Partition:

- Left: (empty)
- Middle: 6 (pivot)
- Right: (empty)

Single element subarray, done, return.

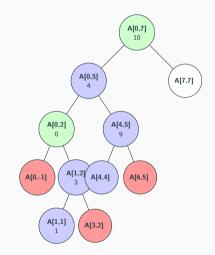


## Randomized Quicksort: Step 3.2.1.2 (Recurse Right [A[6,5]], Done)

Recurse on the right subarray A[6,5] (empty, done).

Return to parent call A[4,5]

Return to parent call A[0,5]



# Randomized Quicksort: Step 3.3 (Recurse Right [A[7,7]], Done)

Recurse on the right subarray A[7,7] After partitioning the right subarray:



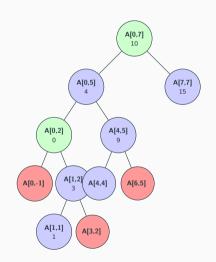
### Partition:

• Left: (empty)

Middle: 1 (pivot)

Right: (empty)

Single element subarray, done, return.



# Randomized Quicksort: Step 4 (Final Sorted Array)

The final sorted array is:

0	1	3	4	6	9	10	15
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# **Quicksort Time Complexity**

• Worst-case:  $O(n^2)$ 

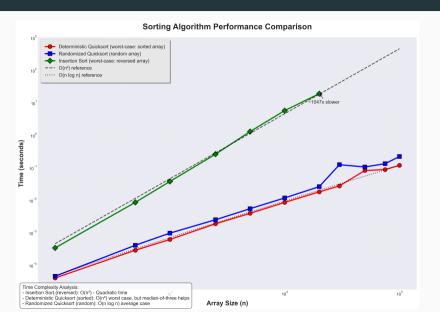
• Best-case:  $O(n \log n)$ 

• Expected:  $O(n \log n)$  (randomized)

**Time Complexity Analysis of** 

Randomized Quicksort

#### **Sorting Algorithms Benchmark**



#### **Quicksort Recurrence**

#### **Expected Comparisons**

Let T(n) be the expected number of comparisons to sort n distinct elements using randomized quicksort:

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

- n comparisons in partitioning: each element compared to the pivot.
- Pivot is chosen uniformly at random.
- For pivot at position i, recursive calls sort subarrays of size i-1 and n-i.
- ullet We average over all n possible pivot positions.

#### **Base Case**

$$T(1) = 0$$
 (single element requires no comparisons)

# Solving the Recurrence Step-by-Step

#### Step 1: Write the Recurrence

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

Step 2: Guess the Solution Assume  $T(n) \le cn \log n$  for some constant c.

# Solving the Recurrence Step-by-Step

#### Step 3: Plug the Guess

$$T(n) \le n + \frac{2c}{n} \sum_{k=1}^{n-1} k \log k$$

Use:

$$\sum_{k=1}^{n-1} k \log k \leq \frac{1}{2} n^2 \log n - \frac{1}{8} n^2$$

Then:

$$T(n) \le n + cn \log n$$

## Step 4: Conclusion

$$T(n) = O(n \log n)$$

# **Example: Monte Carlo Estimation** of pi

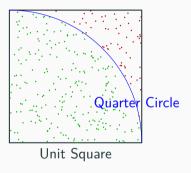
## Monte Carlo Method - Estimating $\pi$

The Monte Carlo method (Metropolis & and, 1949) estimates  $\pi$  by simulating random points in a unit square and counting how many fall inside a quarter circle of radius 1. The ratio of points inside the circle to the total points, multiplied by 4, approximates  $\pi$  (Beckmann, 1971).

#### Monte Carlo Algorithm

- 1. Generate N random points (x,y) where  $0 \le x \le 1$  and  $0 \le y \le 1$ .
- 2. For each point, check if it lies inside the quarter circle:  $x^2+y^2\leq 1$ .
- 3. Count the number of points  ${\cal M}$  that satisfy the condition.
- 4. Estimate  $\pi$  as:  $\pi \approx 4 \times \frac{M}{N}$  (Hammersley & Handscomb, 1964).

#### **Visual Illustration**



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<sup>&</sup>lt;sup>1</sup>This slide was generated using a probabilistic algorithm

## **Example Calculation**

- Suppose we generate N=1000 random points in the unit square
- After simulation, we count  ${\cal M}=785$  points inside the quarter circle
- We estimate  $\pi$  as:

$$\pi \approx 4 \times \frac{M}{N} = 4 \times \frac{785}{1000} = 3.14$$

■ The true value of  $\pi$  is approximately 3.14159 (Beckmann, 1971)

## **Convergence and Error Analysis**

- The error in our estimate decreases as  $O(1/\sqrt{N})$  (Kalos & Whitlock, 2008)
- This means:
  - N=100 points gives roughly 10% error
  - N=10,000 points gives roughly 1% error
  - N=1,000,000 points gives roughly 0.1% error
- The Monte Carlo method is especially useful for calculating multidimensional integrals (Cookson, 2005)
- For  $\pi$  calculation, there are more efficient methods, but this one is visually intuitive

#### **Demo Visualization**



Open Interactive Demo

**Probabilistic Data Structures** 

#### Deterministic vs. Probabilistic Data Structures

# Deterministic (e.g., Hash Set, List):

- Always provide exact answers.
- Can be space-intensive (store all elements).
- Operations might be slower for large datasets (e.g., disk I/O).
- Guarantee: No errors (false positives or negatives).

## Probabilistic (e.g., Bloom Filter):

- Provide approximate answers with controlled error.
- Very space-efficient (use bits, not full elements).
- Operations are typically very fast (constant time).
- Trade-off: Small error probability for huge efficiency gains.

# Key Idea

Use PDS when approximate answers are acceptable and space/speed are critical.

# **Example: Why PDS? Username Availability**

#### The Problem

A website with millions of users needs to instantly check if a username is available during registration. How? (Bloom, 1970)

# Deterministic Approach (Database Query):

- Store all usernames in a database.
- Query DB: 'SELECT 1 FROM users WHERE username = ;
- Accurate? Yes.
- Fast? No. Requires disk I/O, network latency.
- Scalable? Poorly. High load on DB servers.

# Probabilistic Approach (Bloom Filter):

- Keep a compact Bloom filter in memory (Bloom, 1970).
- Check filter: Is 'username' possibly present?
- Accurate? Mostly. Small chance of false positive (saying taken when available), needs DB check then.
- Fast? Yes. In-memory check is O(k).
- Scalable? Excellently. Drastically

reduces DR load

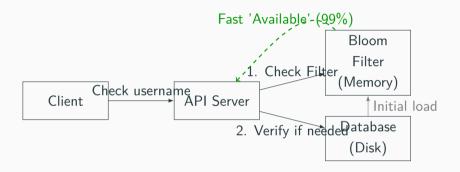
# **Username Checking: Implementation Details**

- 1. **Initialization:** Load all existing usernames into Bloom filter at service startup (only infrequent DB reads).
- 2. New registrations: Add username to both database and Bloom filter.
- 3. Availability check process:
  - Check username against Bloom filter first (Fast, in-memory)
  - If Bloom filter says "definitely not in set"  $\rightarrow$  Username is available (99% case for 1% error rate)
  - If Bloom filter says "possibly in set"  $\rightarrow$  Verify with database query (Slow, but rare)

#### Performance Impact (10M users, 1% error)

- Memory:  $\approx 18$  MB Bloom Filter vs. hundreds of MB for DB index/cache.
- Speed: 99% of availability checks avoid slow database queries. (Wang & Reiter, 2012)

# **Username Checking: System Architecture**



- Bloom filter acts as a fast, efficient preliminary check.
- Deterministic check (DB) used only as a fallback.
- Massively reduces load on the expensive resource (Database).

## **Bloom Filters: The Theory**

- Space-efficient probabilistic data structure (Bloom, 1970)
- Tests if an element is a member of a set
- Possible false positives, never false negatives
- Components:
  - Bit array of m bits (initially all 0)
  - *k* independent hash functions



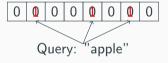
#### **Bloom Filter Operations**

#### Add element:

- 1. Hash element with k functions
- 2. Set bits at these k positions to 1

#### Query element:

- 1. Hash element with k functions
- 2. Check bits at these k positions
- 3. If any bit is 0: Definitely not in set
- 4. If all bits are 1: Probably in set



## The Math Behind Bloom Filters

False positive probability (p):

$$p \approx \left(1 - e^{-kn/m}\right)^k \tag{1}$$

(Broder & Mitzenmacher, 2003)

• Optimal size (m bits) for n items, error p:

$$m = -\frac{n\ln p}{(\ln 2)^2} \tag{2}$$

• Optimal hash functions (*k*):

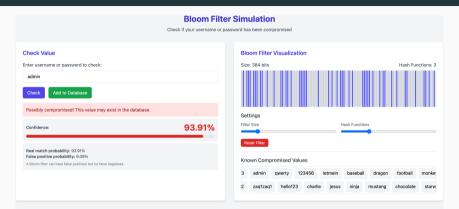
$$k = \frac{m}{n} \ln 2 \approx 0.7 \cdot \frac{m}{n} \tag{3}$$

# **Time and Space Complexity**

Structure	Space	Lookup	Insert	Error Type
Hash Set	O(n)	${\cal O}(1)$ avg	${\cal O}(1)$ avg	None
Bloom Filter	O(m)	O(k)	O(k)	False Positives
Sorted List	O(n)	$O(\log n)$	O(n)	None
Trie	O(N)	O(L)	O(L)	None

n=items, m=bits ( $m \ll n \times item\_size$ ), k=hashes, N=total chars, L=key length

#### **Bloom Filter Simulation**



#### **About Bloom Filters**

A Bloom filter is a space-efficient probabilistic data structure designed to quickly test whether an element is present in a set. It can have false positives (incorrectly reporting an element is in the set) but no false negatives (if it reports an element is not in the set, it definitely is not).

#### How It Works

- 1. Multiple hash functions are applied to the input element
- 2. Each hash function gives a position in the bit array
- 3. When adding an element, all corresponding bits are set to 1
- When checking, if all corresponding bits are 1, the element might be in the set
- 5. If any bit is 0, the element is definitely not in the set

#### Use Cases

- · Checking if a username is taken before querying a database
- · "Have I Been Pwned" password checking
- Spell checkers
- Web cache sharing
- · Network packet routing

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# Other Applications of Bloom Filters

# Web/Database:

- Cache hit/miss optimization (e.g., CDNs)
- Avoid unnecessary DB lookups (like username example)
- Recommendation systems (seen items)
   (Broder & Mitzenmacher, 2003)

#### Network:

- Web crawler URL deduplication (avoid re-crawling)
- Network packet routing (track flows efficiently)
- P2P network resource discovery

# Security:

- Malware signature detection
- Spam filtering (known bad IPs/domains)
- Password breach checking (HavelBeenPwned)

#### Big Data:

- Stream deduplication (unique visitors/events)
- Distributed data sync (approximate differences)
- Genomics (k-mer counting)

#### When to Use Bloom Filters

#### Bloom filters are ideal when:

- Memory is a critical constraint (Big Data, embedded systems)
- False positives are acceptable (can be handled by a secondary check)
- False negatives are unacceptable (must find all true positives)
- Elements are expensive to store or compare
- Lookup speed is crucial (real-time systems)
- Deletions are not needed (or use variants like Counting Bloom Filters)

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