

COMPSCI 514: Algorithms for Data Science

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University of Massachusetts Amherst. Fall 2024.

Lecture 8

Summary

Last Class:

- Finish up Bloom Filters and optimization of number of hash functions.
- Start on streaming algorithms.
- Introduce the frequent items problem and its applications.
- Start on the Count-Min sketch algorithm for frequent items.

This Class:

- Analysis of Count-Min sketch .
- Start on distinct items counting problem.

Approximate Frequent Elements

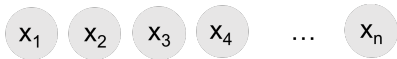
(ϵ, k) -Frequent Items Problem: Consider a stream of n items x_1, \dots, x_n . Return a set F of items, including **all items that appear at least $\frac{n}{k}$ times** and **only items that appear at least $(1 - \epsilon) \cdot \frac{n}{k}$ times**.

- To solve this problem, it suffices to estimate the frequency $f(x)$ of each item x up to error $\pm \frac{\epsilon n}{k}$.
- Will discuss later how to maintain the list of top items in small space.

Count-min sketch:

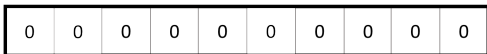
Frequent Elements with Count-Min Sketch

Count-min sketch:



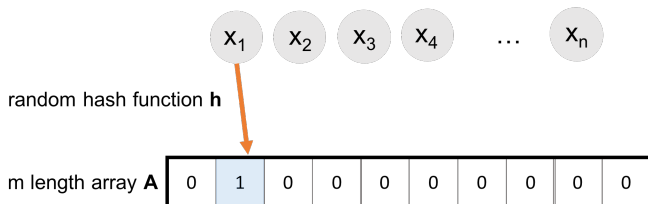
random hash function h

m length array \mathbf{A}



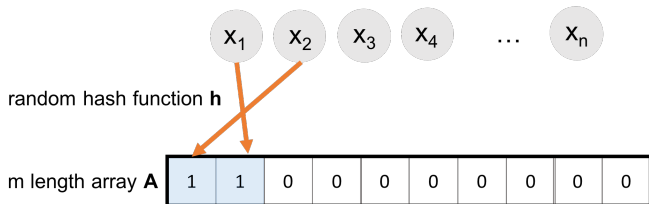
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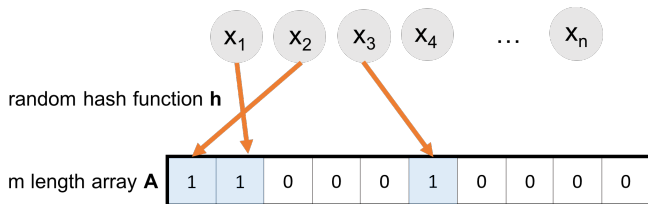
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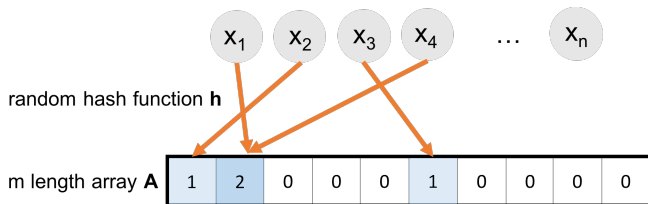
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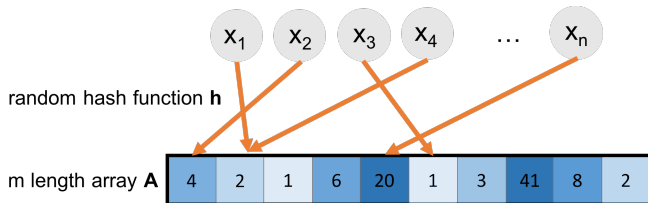
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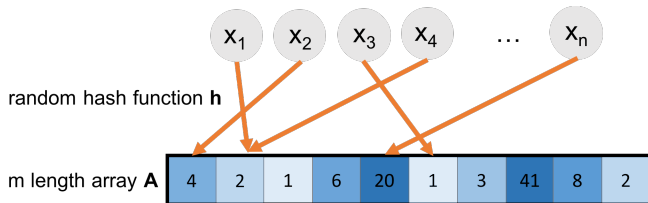
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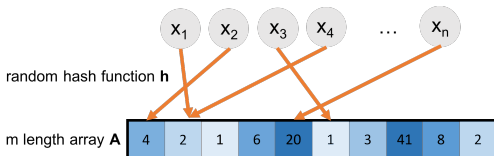
Frequent Elements with Count-Min Sketch

Count-min sketch:



Will use $A[h(x)]$ to estimate $f(x)$, the frequency of x in the stream. I.e., $|\{x_i : x_i = x\}|$.

Count-Min Sketch Accuracy



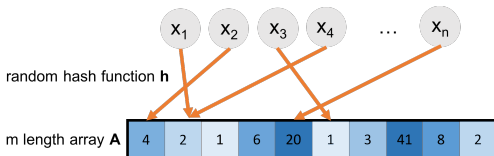
Use $A[h(x)]$ to estimate $f(x)$.

Claim 1: We always have $A[h(x)] \geq f(x)$.

- $A[h(x)]$ counts the number of occurrences of any y with $h(y) = h(x)$, including x itself.

$f(x)$: frequency of x in the stream (i.e., number of items equal to x). h : random hash function. m : size of Count-min sketch array.

Count-Min Sketch Accuracy



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- $A[h(x)]$ counts the number of occurrences of any y with $h(y) = h(x)$, including x itself.
- $A[h(x)] = f(x) + \sum_{y \neq x: h(y)=h(x)} f(y)$.

$f(x)$: frequency of x in the stream (i.e., number of items equal to x). h : random hash function. m : size of Count-min sketch array.

Count-Min Sketch Accuracy

$$A[h(x)] = f(x) + \underbrace{\sum_{y \neq x: h(y) = h(x)} f(y)}_{\text{error in frequency estimate}} .$$

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Expected Error:

$$\mathbb{E} \left[\sum_{y \neq x: h(y)=h(x)} f(y) \right] =$$

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What is a bound on probability that the error is $\geq \frac{2n}{m}$?

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Markov's inequality: $\Pr \left[\sum_{y \neq x: \mathbf{h}(y) = \mathbf{h}(x)} f(y) \geq \frac{2n}{m} \right] \leq \frac{1}{2}$.

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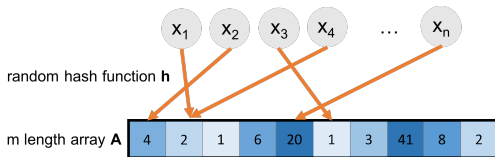
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Markov's inequality: $\Pr \left[\sum_{y \neq x: h(y)=h(x)} f(y) \geq \frac{2n}{m} \right] \leq \frac{1}{2}$.

What property of h is required to show this bound? a) fully random
b) pairwise independent c) 2-universal d) locality sensitive

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Count-Min Sketch Accuracy

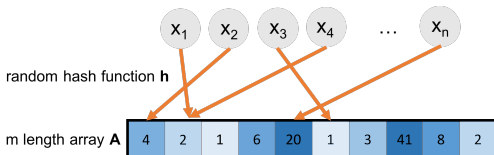


Claim: For any x , with probability at least $1/2$,

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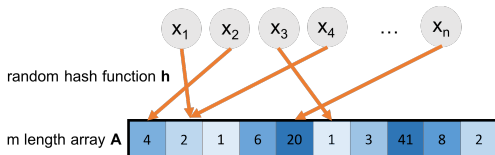
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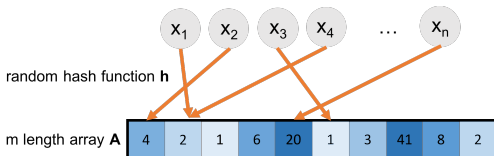
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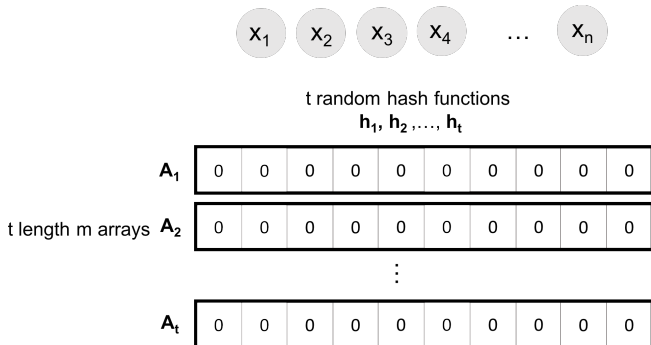
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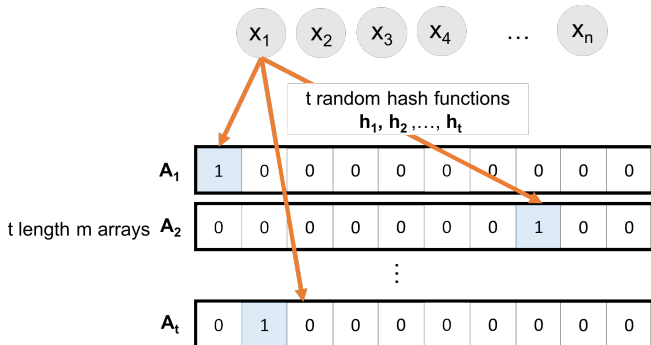
To solve the (ϵ, k) -Frequent elements problem, set $m = \frac{2k}{\epsilon}$. How can we improve the success probability? **Repetition.**

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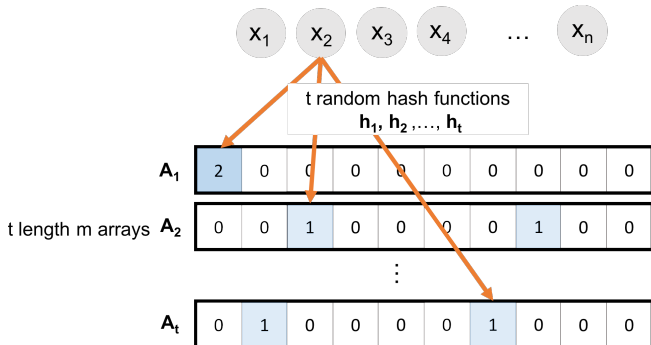
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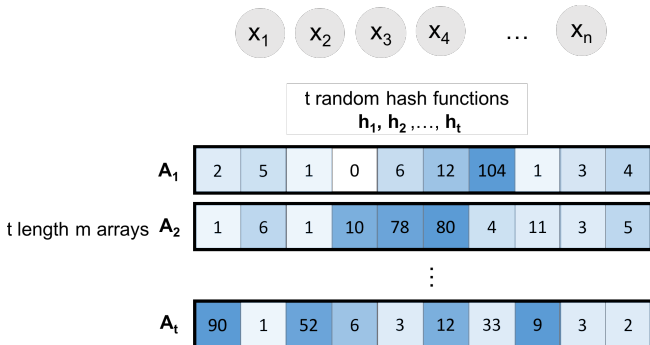
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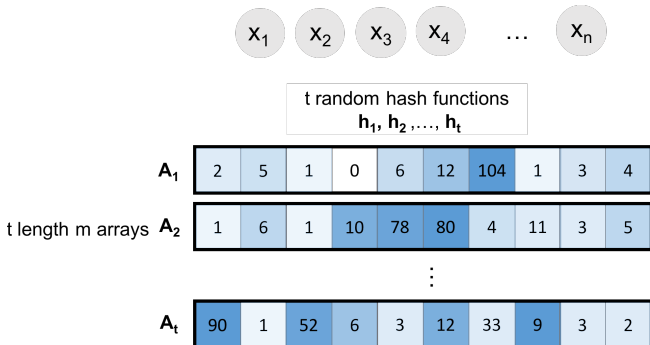
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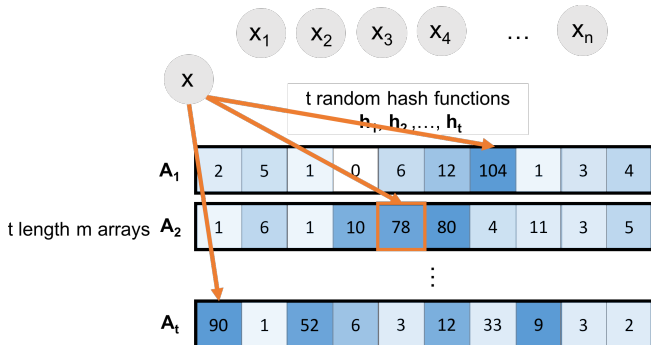


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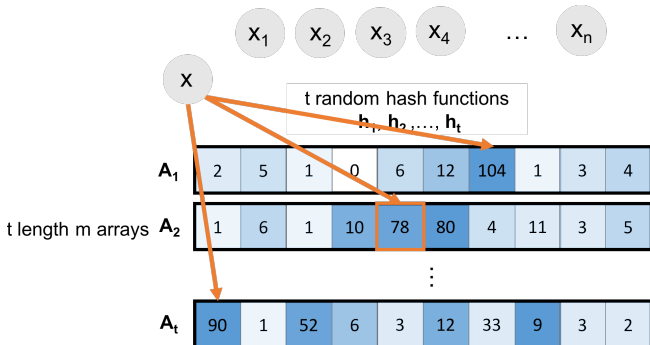
Estimate $f(x)$ with $\tilde{f}(x) = \min_{i \in [t]} A_i[h_i(x)]$. (Count-min sketch)

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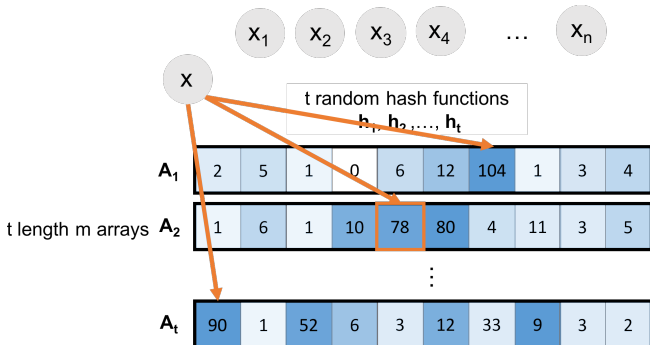
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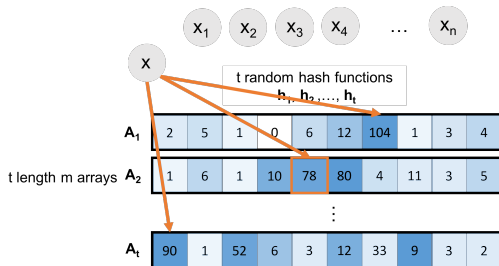
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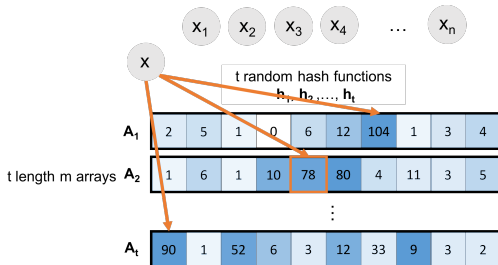
Why min instead of taking the average? The minimum estimate is always the most accurate since they are all overestimates of the true frequency!

Count-Min Sketch Analysis



Estimate $f(x)$ by $\tilde{f}(x) = \min_{i \in [t]} A_i[h_i(x)]$

Count-Min Sketch Analysis

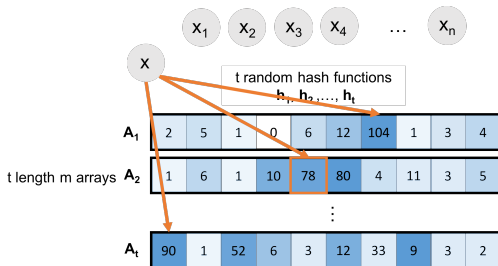


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- For every x and $i \in [t]$, we know that for $m = \frac{2k}{\epsilon}$, with probability $\geq 1/2$:

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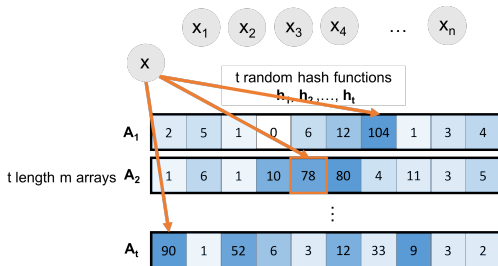
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Count-Min Sketch Analysis



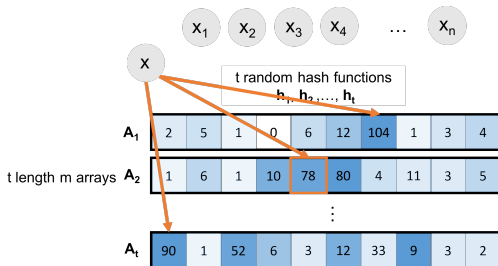
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- What is $\Pr[f(x) \leq \tilde{f}(x) \leq f(x) + \frac{\epsilon n}{k}]$? $1 - 1/2^t$.
- To get a good estimate with probability $\geq 1 - \delta$, set $t = \log_2(1/\delta)$.

Count-Min Sketch

Upshot: Count-min sketch lets us estimate the frequency of each item in a stream up to error $\frac{\epsilon n}{k}$ with probability $\geq 1 - \delta$ in $O(\log(1/\delta) \cdot k/\epsilon)$ space.

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Upshot: Count-min sketch lets us estimate the frequency of each item in a stream up to error $\frac{\epsilon n}{k}$ with probability $\geq 1 - \delta$ in $O(\log(1/\delta) \cdot k/\epsilon)$ space.

- Accurate enough to solve the (ϵ, k) -Frequent elements problem – distinguish between items with frequency $\frac{n}{k}$ and those with frequency $(1 - \epsilon)\frac{n}{k}$.

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Upshot: Count-min sketch lets us estimate the frequency of each item in a stream up to error $\frac{\epsilon n}{k}$ with probability $\geq 1 - \delta$ in $O(\log(1/\delta) \cdot k/\epsilon)$ space.

- Accurate enough to solve the (ϵ, k) -Frequent elements problem – distinguish between items with frequency $\frac{n}{k}$ and those with frequency $(1 - \epsilon)\frac{n}{k}$.
- How should we set δ if we want a good estimate for all items at once, with 99% probability?

Identifying Frequent Elements

Count-min sketch gives an accurate frequency estimate for every item in the stream. But how do we identify the frequent items without having to store/look up the estimated frequency for all elements in the stream?

Identifying Frequent Elements

Count-min sketch gives an accurate frequency estimate for every item in the stream. But how do we identify the frequent items without having to store/look up the estimated frequency for all elements in the stream?

One approach:

- When a new item comes in at step i , check if its estimated frequency is $\geq i/k$ and store it if so.
- At step i remove any stored items whose estimated frequency drops below i/k .
- Store at most $O(k)$ items at once and have all items with frequency $\geq n/k$ stored at the end of the stream.

Questions on Frequent Items?

Distinct Elements

Distinct Elements (Count-Distinct) Problem: Given a stream x_1, \dots, x_n , estimate the number of distinct elements in the stream.

E.g.,

1, 5, 7, 5, 2, 1 \rightarrow 4 distinct elements

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Applications:

- Distinct IP addresses clicking on an ad or visiting a site.
- Distinct values in a database column (for estimating sizes of joins and group bys).
- Number of distinct search engine queries.
- Counting distinct motifs in large DNA sequences.

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Google Sawzall, Facebook Presto, Apache Drill, Twitter Algebird

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Min-Hashing for Distinct Elements (variant of Flajolet-Martin):

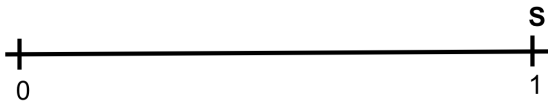
- Let $h : U \rightarrow [0, 1]$ be a random hash function (with a real valued output)
- $s := 1$
- For $i = 1, \dots, n$
 - $s := \min(s, h(x_i))$
- Return $\tilde{d} = \frac{1}{s} - 1$

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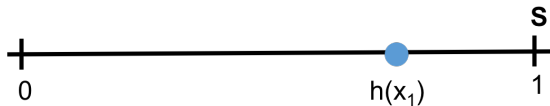


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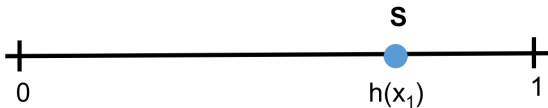


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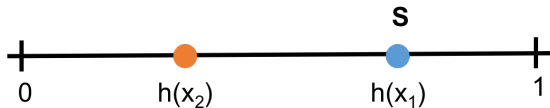


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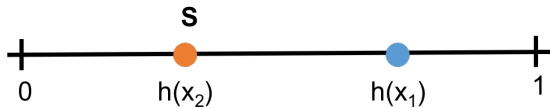


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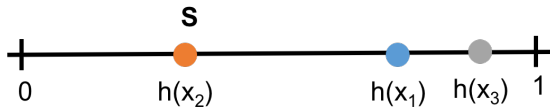


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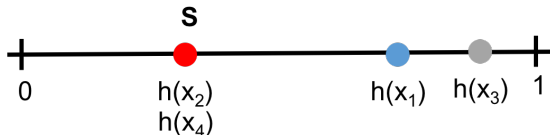


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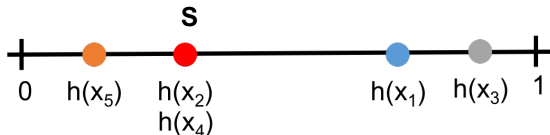


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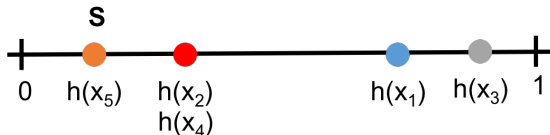


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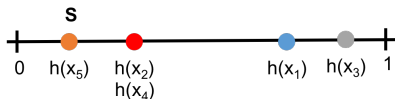
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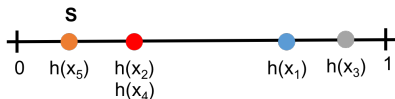
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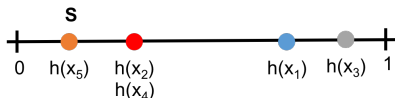


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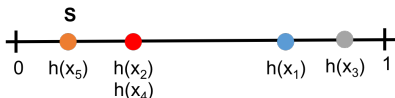


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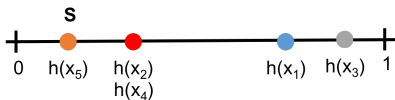
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- Same idea as [Flajolet-Martin algorithm](#) and [HyperLogLog](#), except they use discrete hash functions.

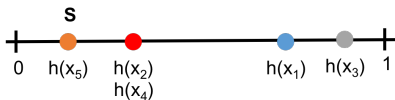
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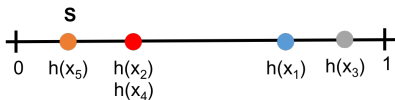
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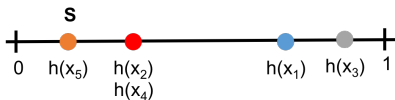
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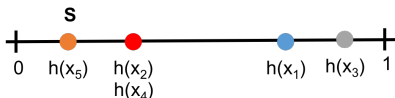


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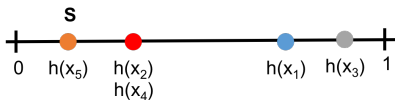


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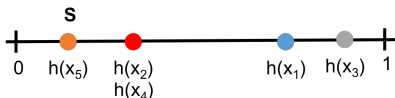


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- **Approximation is robust:** if $|s - \mathbb{E}[s]| \leq \epsilon \cdot \mathbb{E}[s]$ for any $\epsilon \in (0, 1/2)$ and a small constant $c \leq 4$:

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So question is how well \mathbf{s} concentrates around its mean.

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$\mathbf{s} = \frac{1}{k} \sum_{j=1}^k \mathbf{s}_j$. Have already shown that for $j = 1, \dots, k$:

$$\mathbb{E}[\mathbf{s}_j] = \frac{1}{d+1} \implies \mathbb{E}[\mathbf{s}] = \frac{1}{d+1} \text{ (linearity of expectation)}$$

$$\text{Var}[\mathbf{s}_j] \leq \frac{1}{(d+1)^2} \implies \text{Var}[\mathbf{s}] \leq \frac{1}{k \cdot (d+1)^2} \text{ (linearity of variance)}$$

Chebyshev Inequality:

$$\Pr[|\mathbf{s} - \mathbb{E}[\mathbf{s}]| \geq \epsilon \mathbb{E}[\mathbf{s}]] \leq \frac{\text{Var}[\mathbf{s}]}{(\epsilon \mathbb{E}[\mathbf{s}])^2} = \frac{\mathbb{E}[\mathbf{s}]^2/k}{\epsilon^2 \mathbb{E}[\mathbf{s}]^2} = \frac{1}{k \cdot \epsilon^2} = \frac{\epsilon^2 \cdot \delta}{\epsilon^2} = \delta.$$

How should we set k if we want an error with probability at most δ ?

$$k = \frac{1}{\epsilon^2 \cdot \delta}.$$

\mathbf{s}_j : minimum of d distinct hashes chosen randomly over $[0, 1]$. $\mathbf{s} = \frac{1}{k} \sum_{j=1}^k \mathbf{s}_j$.
 $\hat{d} = \frac{1}{\mathbf{s}} - 1$: estimate of # distinct elements d .

Space Complexity

Hashing for Distinct Elements:

- Let $h_1, h_2, \dots, h_k : U \rightarrow [0, 1]$ be random hash functions
- $s_1, s_2, \dots, s_k := 1$
- For $i = 1, \dots, n$
 - For $j=1, \dots, k$, $s_j := \min(s_j, h_j(x_i))$
- $s := \frac{1}{k} \sum_{j=1}^k s_j$
- Return $\hat{d} = \frac{1}{s} - 1$



- Setting $k = \frac{1}{\epsilon^2 \cdot \delta}$, algorithm returns \hat{d} with $|d - \hat{d}| \leq 4\epsilon \cdot d$ with probability at least $1 - \delta$.

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- $\delta = 5\%$ failure rate gives a factor 20 overhead in space complexity.

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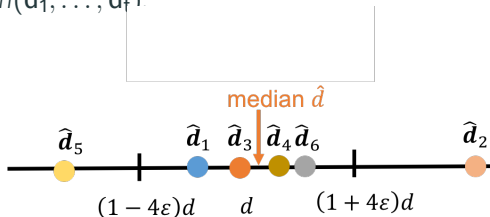
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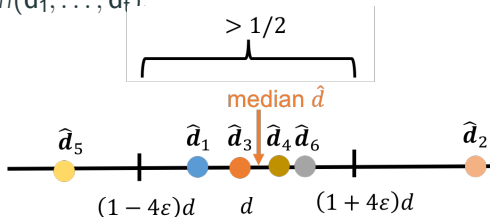


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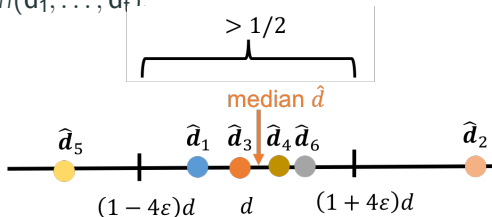
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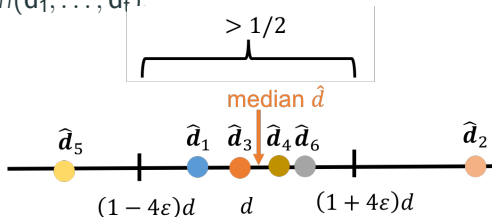
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- If $> 2/3$ of trials fall in $[(1-4\epsilon)d, (1+4\epsilon)d]$, then the median will.
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- $\hat{\mathbf{d}}_1, \dots, \hat{\mathbf{d}}_t$ are the outcomes of the t trials, each falling in $[(1 - 4\epsilon)d, (1 + 4\epsilon)d]$ with probability at least $4/5$.
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Upshot: The median of $t = O(\log(1/\delta))$ independent runs of the hashing algorithm for distinct elements returns $\hat{d} \in [(1 - 4\epsilon)d, (1 + 4\epsilon)d]$ with probability at least $1 - \delta$.

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A note on the median: The median is often used as a robust alternative to the mean, when there are outliers (e.g., heavy tailed distributions, corrupted data).