Count-Min Sketches
Randomization

- Randomization opens up new routes for tradeoffs in data structures:
  - Trade worst-case guarantees for average-case guarantees.
  - Trade exact answers for approximate answers.
- These data structures are used *extensively* in practice. Each of the next lectures is on something you’re likely to encounter IRL.
- Each of the next lectures explores powerful techniques that are useful in navigating the rivers of Theoryland.
Outline for Today

- **Hash Functions**
  - Understanding our basic building blocks.

- **Count-Min Sketches**
  - Estimating how many times we’ve seen something.

- **Concentration Inequalities**
  - “Correct on expectation” versus “correct with high probability.”

- **Probability Amplification**
  - Increasing our confidence in our answers.
Preliminaries: *Hash Functions*
Hashing in Practice

- **Hash functions are used extensively in programming and software engineering:**
  - They make hash tables possible: think C++ `std::hash`, Python’s `__hash__`, or Java’s `Object.hashCode()`.
  - They’re used in cryptography: SHA-256, HMAC, etc.
- **Question:** When we’re in Theoryland, what do we mean when we say “hash function?”
Hashing in Theoryland

- In Theoryland, a hash function is a function from some domain called the *universe* (typically denoted $\mathcal{U}$) to some codomain.
- The codomain is usually a set of the form $[m] = \{0, 1, 2, 3, \ldots, m - 1\}$

$$h : \mathcal{U} \rightarrow [m]$$
Hashing in Theoryland

- **Intuition:** No matter how clever you are with designing a hash function, that hash function isn’t random, and so there will be pathological inputs.
  - You can formalize this with the pigeonhole principle.
- **Idea:** Rather than finding the One True Hash Function, we’ll assume we have a collection of hash functions to pick from, and we’ll choose which one to use randomly.
Families of Hash Functions

- A **family** of hash functions is a set $\mathcal{H}$ of hash functions with the same domain and codomain.
- We can then introduce randomness into our data structures by sampling a random hash function from $\mathcal{H}$.
- **Key Point:** The randomness in our data structures almost always derives from the random choice of hash functions, not from the data.
  
  *Data is adversarial.*
  
  *Hash function selection is random.*

- **Question:** What makes a family of hash functions $\mathcal{H}$ a “good” family of hash functions?
**Goal:** If we pick $h \in \mathcal{H}$ uniformly at random, then $h$ should distribute elements uniformly randomly.

**Problem:** A hash function that distributes $n$ elements uniformly at random over $[m]$ requires $\Omega(n \log m)$ space in the worst case.

**Question:** Do we actually need true randomness? Or can we get away with something weaker?
**Distribution Property:**
Each element should have an equal probability of being placed in each slot.

For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

Some “obviously bad” hash functions obey this rule. How is this possible?
**Distribution Property:**
Each element should have an equal probability of being placed in each slot.

For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

**Problem:** This rule doesn’t guarantee that elements are spread out.
**Distribution Property:** Each element should have an equal probability of being placed in each slot.

**Independence Property:** Where one element is placed shouldn’t impact where a second goes.

For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

For any distinct $x, y \in \mathcal{U}$ and random $h \in \mathcal{H}$, $h(x)$ and $h(y)$ are independent random variables.
**Distribution Property:**
Each element should have an equal probability of being placed in each slot.

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Where one element is placed shouldn’t impact where a second goes.

For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

For any distinct $x, y \in \mathcal{U}$ and random $h \in \mathcal{H}$, $h(x)$ and $h(y)$ are independent random variables.

A family of hash functions $\mathcal{H}$ is called **2-independent** (or **pairwise independent**) if it satisfies the distribution and independence properties.
For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$. 

**Intuition:**
2-independence means any pair of elements is unlikely to collide.

For any distinct $x, y \in \mathcal{U}$ and random $h \in \mathcal{H}$, $h(x)$ and $h(y)$ are independent random variables.

\[
\Pr[h(x) = h(y)] = \sum_{i=0}^{m-1} \Pr[h(x) = i] \cdot \Pr[h(y) = i]
\]
For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

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\Pr[h(x) = h(y)] = \sum_{i=0}^{m-1} \Pr[h(x) = i \land h(y) = i] = \sum_{i=0}^{m-1} \Pr[h(x) = i] \cdot \Pr[h(y) = i] = \sum_{i=0}^{m-1} \frac{1}{m^2}
\]
For any $x \in \mathcal{U}$ and random $h \in \mathcal{H}$, the value of $h(x)$ is uniform over $[m]$.

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= \sum_{i=0}^{m-1} \frac{1}{m^2}
= \frac{1}{m}
$$

This is the same as if $h$ were a truly random function.

For any distinct $x, y \in \mathcal{U}$ and random $h \in \mathcal{H}$, $h(x)$ and $h(y)$ are independent random variables.
For more on hashing outside of Theoryland, check out this Stack Exchange post.
Approximating Quantities
What makes for a good “approximate” solution?
What does it mean for an approximation to be “good”?

Let \( A \) be the true answer. Let \( \hat{A} \) be a random variable denoting our estimate.

This would not make for a good estimate. However, we have \( E[\hat{A}] = A \).

**Observation 1:** Being correct in expectation isn’t sufficient.
Let $\mathbf{A}$ be the true answer. Let $\hat{\mathbf{A}}$ be a random variable denoting our estimate.

It’s unlikely that we’ll get the right answer, but we’re probably going to be close.

**Observation 2:** The difference $|\hat{\mathbf{A}} - \mathbf{A}|$ between our estimate and the truth should ideally be small.

What does it mean for an approximation to be “good”?
What does it mean for an approximation to be “good”?

Let $A$ be the true answer. Let $Â$ be a random variable denoting our estimate.

This estimate skews low, but it’s very close to the true value.

**Observation 3:** An estimate doesn’t have to be unbiased to be useful.

**Distribution of our estimate $Â$.**

**$A$**

(True answer)
What does it mean for an approximation to be “good”?

Let $A$ be the true answer. Let $\hat{A}$ be a random variable denoting our estimate.

The more resources we allocate, the better our estimate should be.

*Observation 4:* A good approximation should be tunable.
What does it mean for an approximation to be “good”?

Suppose there are two tunable values

\[ \varepsilon \in (0, 1] \]
\[ \delta \in (0, 1] \]

where \( \varepsilon \) represents \textit{accuracy} and \( \delta \) represents \textit{confidence}.

\textbf{Goal:} Make an estimator \( \hat{A} \) for some quantity \( A \) where

\[
\text{With probability at least } 1 - \delta, \quad |\hat{A} - A| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of the size of the input.
**Goal:** Make an estimator \( \hat{A} \) for some quantity \( A \) where

\[
\text{With probability at least } 1 - \delta, \quad |A - \hat{A}| \leq \epsilon \cdot \text{size(input)}
\]

for some measure of the size of the input.

\( \delta = \frac{1}{2} \)

\( \epsilon \) small
What does it mean for an approximation to be “good”?

Goal: Make an estimator \( \hat{A} \) for some quantity \( A \) where

\[
|A - \hat{A}| \leq \epsilon \cdot \text{size}(\text{input})
\]

for some measure of the size of the input.

\( \delta = \frac{1}{2} \)
\( \epsilon \) medium

Correct

True answer
What does it mean for an approximation to be “good”?

**Goal:** Make an estimator \( \hat{A} \) for some quantity \( A \) where

\[
\text{With probability at least } 1 - \delta, \quad |A - \hat{A}| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of the size of the input.

\( \delta = \frac{1}{2} \)

\( \varepsilon \) large

Correct

True answer
What does it mean for an approximation to be “good”?

Goal: Make an estimator $\hat{A}$ for some quantity $A$ where

\[
\text{With probability at least } 1 - \delta, \quad |A - \hat{A}| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of the size of the input.

$\delta = \frac{1}{16}$

$\varepsilon$ small

Correct

True answer

Probably

Approximately
Frequency Estimation
Frequency Estimators

• A frequency estimator is a data structure supporting the following operations:
  • $\text{increment}(x)$, which increments the number of times that $x$ has been seen, and
  • $\text{estimate}(x)$, which returns an estimate of the frequency of $x$.

• Using BSTs, we can solve this in space $\Theta(n)$ with worst-case $O(\log n)$ costs on the operations.

• Using hash tables, we can solve this in space $\Theta(n)$ with expected $O(1)$ costs on the operations.
Frequency Estimators

- Frequency estimation has many applications:
  - Search engines: Finding frequent search queries.
  - Network routing: Finding common source and destination addresses.
- In these applications, $\Theta(n)$ memory can be impractical.
- **Goal:** Get approximate answers to these queries in sublinear space.
The Count-Min Sketch
How to Build an Estimator

1. Design a simple data structure that, intuitively, gives you a good estimate.

2. Use a *sum of indicator variables* and *linearity of expectation* to prove that, on expectation, the data structure is pretty close to correct.

3. Use a *concentration inequality* to show that, with decent probability, the data structure’s output is close to its expectation.

4. Run multiple copies of the data structure in parallel to amplify the success probability.
Revisiting the Exact Solution

- In the exact solution to the frequency estimation problem, we maintained a single counter for each distinct element. This is too space-inefficient.

- **Idea:** Store a fixed number of counters and assign a counter to each $x_i \in \mathcal{U}$. Multiple $x_i$'s might be assigned to the same counter.

- To *increment*($x$), increment the counter for $x$.

- To *estimate*($x$), read the value of the counter for $x$. 

![Diagram of different counters with numbers 11, 6, 4, and 7]
Our Initial Structure

- We can model “assigning each $x_i$ to a counter” by using hash functions.
- Choose, from a family of 2-independent hash functions $\mathcal{H}$, a uniformly-random hash function $h : \mathcal{U} \rightarrow [w]$.
- Create an array $\text{count}$ of $w$ counters, each initially zero.
  - We'll choose $w$ later on.
- To $\text{increment}(x)$, increment $\text{count}[h(x)]$.
- To $\text{estimate}(x)$, return $\text{count}[h(x)]$.
Analyzing our Structure
For each \( x_i \in \mathcal{U} \), let \( a_i \) denote the number of times we’ve seen \( x_i \).

Similarly, let \( \hat{a}_i \) denote our estimated value of the frequency of \( x_i \).

**Goal:** Bound the probability that the error \((\hat{a}_i - a_i)\) is too high.
**Idea:** Think of our element frequencies $a_1, a_2, a_3, \ldots$ as a vector $a = [a_1, a_2, a_3, \ldots]$. The total number of objects is the sum of the vector entries. This is called the $L_1$ norm of $a$, and is denoted $\|a\|_1$: $\|a\|_1 = \sum_i |a_i|$.

There are $\|a\|_1$ total elements distributed across $w$ buckets. We’re using a 2-independent hash family.

**Reasonable guess:** each bin has $\|a\|_1 / w$ elements in it, so $\hat{a}_i - a_i \leq \|a\|_1 / w$.

**Question:** Intuitively, what should we expect our approximation error to be?
Analyzing this Structure

Let's look at \( \hat{a}_i = \text{count}[h(x_i)] \) for some choice of \( x_i \).

For each element \( x_j \):

- If \( h(x_i) = h(x_j) \), then \( x_j \) contributes \( a_j \) to \( \text{count}[h(x_i)] \).
- If \( h(x_i) \neq h(x_j) \), then \( x_j \) contributes 0 to \( \text{count}[h(x_i)] \).

To pin this down precisely, let’s define a set of random variables \( X_1, X_2, \ldots \), as follows:

\[
X_j = \begin{cases} 
1 & \text{if } h(x_i) = h(x_j) \\
0 & \text{otherwise}
\end{cases}
\]

Each of these variables is called an *indicator random variable*, since it “indicates” whether some event occurs.
Analyzing this Structure

• Let's look at $\hat{a}_i = \text{count}[h(x_i)]$ for some choice of $x_i$.
• For each element $x_j$:
  • If $h(x_i) = h(x_j)$, then $x_j$ contributes $a_j$ to $\text{count}[h(x_i)]$.
  • If $h(x_i) \neq h(x_j)$, then $x_j$ contributes 0 to $\text{count}[h(x_i)]$.
• To pin this down precisely, let’s define a set of random variables $X_1, X_2, \ldots$, as follows:

\[ X_j = \begin{cases} 
1 & \text{if } h(x_i) = h(x_j) \\
0 & \text{otherwise}
\end{cases} \]

• The value of $\hat{a}_i - a_i$ is then given by

\[ \hat{a}_i - a_i = \sum_{j \neq i} a_j X_j \]
\[ E[\hat{a}_i - a_i] = E\left[ \sum_{j \neq i} a_j X_j \right] \]

\[ = \sum_{j \neq i} E[a_j X_j] \]

This follows from **linearity of expectation**. We’ll use this property extensively over the next few days.
\[
E[\hat{a}_i - a_i] = E\left[ \sum_{j \neq i} a_j X_j \right] \\
= \sum_{j \neq i} E[ a_j X_j ] \\
= \sum_{j \neq i} a_j E[ X_j ]
\]

The values of \( a_j \) are not random. \textit{The randomness comes from our choice of hash function.}
\begin{align*}
E[\hat{a}_i - a_i] &= E[\sum_{j \neq i} a_j X_j] \\
&= \sum_{j \neq i} E[a_j X_j] \\
&= \sum_{j \neq i} a_j E[X_j]
\end{align*}

\begin{equation*}
E[X_j] = 1 \cdot \Pr[h(x_i) = h(x_j)] + 0 \cdot \Pr[h(x_i) \neq h(x_j)]
\end{equation*}

\begin{equation*}
X_j = \begin{cases} 
1 & \text{if } h(x_i) = h(x_j) \\
0 & \text{otherwise}
\end{cases}
\end{equation*}
\[
E[\hat{a}_i - a_i] = E[\sum_{j \neq i} a_j X_j] = \sum_{j \neq i} E[a_j X_j] = \sum_{j \neq i} a_j E[X_j]
\]

If \( X \) is an indicator variable for some event \( \mathcal{E} \), then \( E[X] = Pr[\mathcal{E}] \). This is really useful when using linearity of expectation!
\[ E[\hat{a}_i - a_i] = E[\sum_{j \neq i} a_j X_j] \]

\[ = \sum_{j \neq i} E[a_j X_j] \]

\[ = \sum_{j \neq i} a_j E[X_j] \]

\[ E[X_j] = 1 \cdot \Pr[h(x_i) = h(x_j)] + 0 \cdot \Pr[h(x_i) \neq h(x_j)] \]

\[ = \Pr[h(x_i) = h(x_j)] \]

\[ = \frac{1}{w} \]

Hey, we saw this earlier!
\[ E[\hat{a}_i - a_i] = E[\sum_{j \neq i} a_j X_j] \]
\[ = \sum_{j \neq i} E[a_j X_j] \]
\[ = \sum_{j \neq i} a_j E[X_j] \]
\[ = \sum_{j \neq i} \frac{a_j}{w} \]
\[ \leq \frac{\|a\|_1}{w} \]

\[ E[X_j] = 1 \cdot \Pr[h(x_i) = h(x_j)] + 0 \cdot \Pr[h(x_i) \neq h(x_j)] \]
\[ = \Pr[h(x_i) = h(x_j)] \]
\[ = \frac{1}{w} \]
Goal: Make an estimator \( \hat{a} \) for some quantity \( a \) where

With probability at least 1 - \( \delta \),

\[
|\hat{a} - a| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of the size of the input.

How do we tune \( w \) so we’re likely to fall in this range?

\[
\mathbb{E}[\hat{a}_i - a_i] \leq \frac{\|a\|_1}{w}
\]
\[ \Pr \left[ \hat{a}_i - a_i > \varepsilon \| a \|_1 \right] \leq \frac{\mathbb{E} \left[ \hat{a}_i - a_i \right]}{\varepsilon \| a \|_1} \]

We don’t know the exact distribution of this random variable.

However, we have a **one-sided error**: our estimate can never be lower than the true value. This means that \( \hat{a}_i - a_i \geq 0 \).

**Markov’s inequality** says that if \( X \) is a nonnegative random variable, then

\[ \Pr[X \geq c] \leq \frac{\mathbb{E}[X]}{c} \]
\[ \text{Pr}[\hat{a}_i - a_i > \varepsilon \|a\|_1] \leq \frac{E[\hat{a}_i - a_i]}{\varepsilon \|a\|_1} \leq \frac{\|a\|_1}{w} \cdot \frac{1}{\varepsilon \|a\|_1} \]

\[ E[\hat{a}_i - a_i] \leq \frac{\|a\|_1}{w} \]
\[ \Pr [ \hat{a}_i - a_i > \varepsilon \|a\|_1 ] \]

\[ \leq \frac{\mathbb{E} [ \hat{a}_i - a_i ]}{\varepsilon \|a\|_1} \]

\[ \leq \frac{\|a\|_1 \cdot \frac{1}{w}}{\varepsilon \|a\|_1} \]

\[ = \frac{1}{\varepsilon w} \]
**Goal:** Make an estimator \( \hat{a} \) for some quantity \( a \) where

With probability at least \( 1 - \delta \),

\[
|\hat{a} - a| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of input size.

\[
\text{Pr}[\hat{a}_i - a_i > \varepsilon \|a\|_1] \leq \frac{1}{\varepsilon w}
\]

**Initial Idea:**
Pick \( w = \varepsilon^{-1} \cdot \delta^{-1} \). Then

\[
\text{Pr}[\hat{a}_i - a_i > \varepsilon \|a\|_1] \leq \delta
\]

**Suppose we’re counting 1,000 distinct items.**

If we want our estimate to be within \( \varepsilon \|a\|_1 \) of the true value with 99.9% probability, how much memory do we need?

**Answer:** \( 1,000 \cdot \varepsilon^{-1} \).

**Can we do better?**
**Goal:** Make an estimator \( \hat{a} \) for some quantity \( a \) where

With probability at least \( 1 - \delta \),

\[
|\hat{a} - a| \leq \varepsilon \cdot \text{size(input)}
\]

for some measure of input size.

\[
\Pr[\hat{a}_i - a_i > \varepsilon \|a\|_1] \leq \frac{1}{\varepsilon w}
\]

**Revised Idea:** Pick \( w = e \cdot \varepsilon^{-1} \). Then

\[
\Pr[\hat{a}_i - a_i > \varepsilon \|a\|_1] < e^{-1}
\]

This simple data structure, by itself, is likely to be wrong.

What happens if we run a bunch of copies of this approach in parallel?
Let's suppose that we run $d$ independent copies of this data structure. Each has its own independently randomly chosen hash function.

To `increment(x)` in the overall structure, we call `increment(x)` on each of the underlying data structures.

The probability that at least one of them provides a good estimate is quite high.

**Question:** How do you know which one?

<table>
<thead>
<tr>
<th>Estimator 1:</th>
<th>Estimator 2:</th>
<th>Estimator 3:</th>
<th>Estimator 4:</th>
<th>Estimator 5:</th>
</tr>
</thead>
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<tr>
<td>137</td>
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<td>166</td>
<td>103</td>
<td>261</td>
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</table>
Running in Parallel

• Let's suppose that we run $d$ independent copies of this data structure. Each has its own independently randomly chosen hash function.

• To $\text{increment}(x)$ in the overall structure, we call $\text{increment}(x)$ on each of the underlying data structures.

• The probability that at least one of them provides a good estimate is quite high.

• **Question:** How do you know which one?

**Intuition:** The smallest estimate returned has the least “noise,” and that’s the best guess for the frequency.

<table>
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</table>
\[
\Pr \left[ \min \{ \hat{a}_{ij} \} - a_i > \varepsilon \|a\|_1 \right]
\]

\[
= \Pr \left[ \bigwedge_{j=1}^{d} \left( \hat{a}_{ij} - a_i > \varepsilon \|a\|_1 \right) \right]
\]

The only way the minimum estimate is inaccurate is if every estimate is inaccurate.

Let \( \hat{a}_{ij} \) be the estimate from the \( j \)th copy of the data structure.

Our final estimate is \( \min \{ \hat{a}_{ij} \} \)
\[
\Pr \left[ \min \{ \hat{a}_{ij} \} - a_i > \epsilon \|a\|_1 \right] = \Pr \left[ \bigwedge_{j=1}^{d} \left( \hat{a}_{ij} - a_i > \epsilon \|a\|_1 \right) \right] = \prod_{j=1}^{d} \Pr \left[ \hat{a}_{ij} - a_i > \epsilon \|a\|_1 \right]
\]

Each copy of the data structure is independent of the others.

Let \(\hat{a}_{ij}\) be the estimate from the \(j\)th copy of the data structure.

Our final estimate is \(\min \{ \hat{a}_{ij} \}\).
Let $\hat{a}_{ij}$ be the estimate from the $j$th copy of the data structure. Our final estimate is $\min \{ \hat{a}_{ij} \}$. 

\[
\Pr \left[ \min \{ \hat{a}_{ij} \} - a_i > \varepsilon \| a \|_1 \right] \\
= \Pr \left[ \bigwedge_{j=1}^{d} (\hat{a}_{ij} - a_i > \varepsilon \| a \|_1) \right] \\
= \prod_{j=1}^{d} \Pr [ \hat{a}_{ij} - a_i > \varepsilon \| a \|_1 ] \\
\leq \prod_{j=1}^{d} e^{-1}
\]
Let $\hat{a}_{ij}$ be the estimate from the $j$th copy of the data structure. Our final estimate is $\min \{ \hat{a}_{ij} \}$.
Goal: Make an estimator $\hat{a}$ for some quantity $a$ where

With probability at least $1 - \delta$,

$$|\hat{a} - a| \leq \varepsilon \cdot \text{size(input)}$$

for some measure of input size.

$$\Pr[\min\{\hat{a}_{ij} - a_i\} > \varepsilon \|a\|_1] \leq e^{-d}$$

Idea: Choose $d = -\ln \delta$. (Equivalently: $d = \ln \delta^{-1}$.) Then

$$\Pr[\min\{\hat{a}_{ij} - a_i\} > \varepsilon \|a\|_1] \leq \delta$$
The Count-Min Sketch

\[ w = \lceil e \cdot \varepsilon^{-1} \rceil \]

\[ d = \lceil \ln \delta^{-1} \rceil \]

Sampled uniformly and independently from a 2-independent family of hash functions
The Count-Min Sketch

increment(x):
    for i = 1 ... d:
        count[i][h_i(x)]++

estimate(x):
    result = ∞
    for i = 1 ... d:
        result = min(result, count[i][h_i(x)])
    return result
The Count-Min Sketch

• Update and query times are $\Theta(d)$, which is $\Theta(\log \delta^{-1})$.

• Space usage: $\Theta(\varepsilon^{-1} \cdot \log \delta^{-1})$ counters.
  • This is a major improvement over our earlier approach that used $\Theta(\varepsilon^{-1} \cdot \delta^{-1})$ counters.
  • This can be significantly better than just storing a raw frequency count!

• Provides an estimate to within $\varepsilon \|a\|_1$ with probability at least $1 - \delta$. 
Major Ideas From Today

- **2-independent hash families** are useful when we want to keep collisions low.
- A “good” approximation of some quantity should have tunable **confidence** and **accuracy** parameters.
- **Sums of indicator variables** are useful for deriving expected values of estimators.
- **Concentration inequalities** like **Markov’s inequality** are useful for showing estimators don’t stay too much from their expected values.
- Good estimators can be built from multiple parallel copies of weaker estimators.
Next Time

- **Count Sketches**
  - An alternative frequency estimator with different time/space bounds.

- **Bloom Filters**
  - Storing data in close to theoretically optimal space.