

## Data Sketching

The Bloom Filter  
*(membership with controlled-error)*

## Dictionary problem

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What data structures you know for storing a  
**set of keys** and supporting **exact searches**  
and **insert operations** over them?

Hashing

What about false positives?

<b>Bloom Filters</b> Count-Min Sketches	<b>The problem</b> Main idea Mathematics Compressed Bloom Filters Spectral Bloom Filters Some applications
<b>Reminder</b>	
<p><i>Wherever a list or set is used, and space is a consideration, a Bloom Filter should be considered.</i></p> <p><i>When using a Bloom Filter, consider the effects of false positives.</i></p>	
Bloom [1970]	

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<b>Membership query</b>	
<b>Definition</b>	
The <i>Membership Problem</i>	
◇ Given a set $S$ and an element $y$ : $y \in S$ ?	
◇ Given a set $S$ compute its characteristic function $\chi_S$	
$\chi_S(y) = \begin{cases} 1, & \text{if } y \in S \\ 0, & \text{if } y \notin S \end{cases}$	
• well-known solutions	
• Linear Scan	
• Hash Functions	

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Preconditions

- given a set of objects  $S = \{x_1, \dots, x_n\}$ 
  - no restrictions on objects
  
- a vector  $B$  of  $m$  bits where  $b_i \in \{0, 1\}$ 
  - we will discuss next about the value of  $m$
  
- suppose we have  $k$  hash functions  $h_1, \dots, h_k$ 
  - each  $h_i$  is defined as  $h_i : U \supseteq S \rightarrow [1; m]$

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Building  $B[1, m]$

- For each  $x \in S$ , we set  $B[h_j(x)] = 1, \forall j = 1, 2, \dots, k$ .

$B$ 

1	0	1	.....	0	1	.....	1	0	
$h_{i_1}(x_i)$		$h_{j_1}(x_j)$		$h_{i_k}(x_i)$			$h_{j_k}(x_j)$		

The *build-time* is  $\Theta(k|S|)$  time and  $\Theta(|B|) = \Theta(m)$  space

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## Searching in $B[1, m]$

- We claim " $y \in S$ " if  $b_{h_i(y)} = 1, \forall i = 1, \dots, k$ .

$y \notin S$

The *search*-time is  $O(k)$  time.

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## The main problem

**Definition**

The main problem are *false positives*:  
it may exist  $x_j \neq y$  such that  $h_i(x_j) = h_i(y), \forall i = 1, \dots, k$ .

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Bloom Filters and CM-Sketches

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### Example: building $B$

	$h_1$	$h_2$	$h_3$
ACG	3	6	4
ATA	2	11	10
CGA	11	9	6
TTA	1	10	9
TTT	6	2	1
CGC	9	3	3
AAA	4	1	2
TCT	10	4	11
...	...	...	...

$S = \{TTA, TCT, \text{ATA}\}$

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### Example: searching $B$

	$h_1$	$h_2$	$h_3$
ACG	3	6	4
ATA	2	11	10
CGA	11	9	6
TTA	1	10	9
TTT	6	2	1
CGC	9	3	3
AAA	4	1	2
TCT	10	4	11
...	...	...	...

$S = \{TTA, TCT, \text{ATA}\}$

$CGA \in S \rightarrow NO$

$B_{h_3}(CGA) = 0$

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## Example: searching $B$

$h_1(TTA) = h_2(AAA) = 1$   
 $h_2(ATA) = h_3(AAA) = 2$   
 $h_2(TCT) = h_1(AAA) = 4$

$S = \{TTA, TCT, ATA\}$

	$h_1$	$h_2$	$h_3$
ACG	3	6	4
ATA	2	11	10
CGA	11	9	6
TTA	1	10	9
TTT	6	2	1
CGC	9	3	3
AAA	4	1	2
TCT	10	4	11
...	...	...	...

$AAA \stackrel{?}{\in} S \rightarrow YES$   
*false positive*

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<b>Probability of a <i>false positive</i></b>	
<ul style="list-style-type: none"><li>• assumption that hash are perfectly random</li><li>• after <i>build</i></li></ul> $\mathcal{P}(b_i = 0) = \left(1 - \frac{1}{m}\right)^{kn} \approx e^{-kn/m} = p$ <ul style="list-style-type: none"><li>• probability of a <i>false positive</i> is</li></ul> $(1 - e^{-kn/m})^k = (1 - p)^k = \varepsilon$ <div style="text-align: right; background-color: red; color: white; padding: 2px;">Not perfectly true but...</div> <ul style="list-style-type: none"><li>• other formulations are <i>asymptotically</i> equivalent</li></ul>	

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<b>Optimizing number of <i>hash functions</i></b>	
<ul style="list-style-type: none"><li>• higher <i>k</i>-value: more chances to find a 0-bit for <math>y \notin S</math>.</li><li>• lower <i>k</i>-value: increase fraction of 0-bits in <i>B</i>.</li><li>• minimize the <math>\varepsilon</math> function</li></ul> $\tilde{k} = \ln 2 \cdot (m/n)$ <ul style="list-style-type: none"><li>• With this value of <math>\tilde{k}</math>, we have <math>p = 0.5</math> and thus</li></ul> $\varepsilon = (0.5)^{\tilde{k}} = (0.6185)^{m/n}$	

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## How big should be the $B$ vector?

- depends on the  $\varepsilon$  value we want: given  $n$  we fix  $m$ .

$m$	$\varepsilon$
$n$	0.61
$2n$	0.38
$5n$	0.09
$10n$	0.008

$m = \Theta(n)$  is generally a good choice

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## Bloom Filters v.s. hash functions

$\diamond$	hash functions	Bloom Filters
<i>build time</i>	$\Theta(n)$	$\Theta(n)$
<i>space needed</i>	$\Theta(n \log n)$	$\Theta(m)$
<i>search time</i>	$O(1)$	$(m/n) \ln 2$
$\varepsilon$ value	.	$(0.6185)^{m/n}$

- Hash functions are Bloom Filters with  $k = 1$



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<b>Bloom Filters tricks</b>	
<ul style="list-style-type: none"><li>• union by <i>OR</i><ol style="list-style-type: none"><li>1 we have sets <math>S_1, S_2</math> and Bloom Filters <math>B^1, B^2</math></li><li>2 suppose <math>m_1 = m_2</math> and same hashing functions</li><li>3 just OR the counters</li></ol><math display="block">B_i^{12} = B_i^1 \vee B_i^2</math></li> <li>• halved size<ol style="list-style-type: none"><li>1 suppose <math>m = 2^\alpha</math></li><li>2 make union by <i>OR</i> of the two halves</li><li>3 when hashing, mask high-order bit</li></ol></li></ul>	

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<b>Spectral Bloom Filters (SBF)</b>	
<div style="border: 1px solid black; padding: 5px;"><p><b>Definition</b></p><p><math>M = \langle S, f_x \rangle</math> is a multiset where</p><ul style="list-style-type: none"><li>• <math>S</math> is a set</li><li>• <math>f_x</math> is a function returning the #occurrences of <math>x</math> in <math>M</math></li></ul></div> <p>Notice that a <i>stream</i> might be looked at as a <i>multiset</i>.</p> <p><u>ex</u> Given <math>\{A, A, B, C, C\}</math> We have <math>S = \{A, B, C\}</math> and <math>f_A = f_C = 2, f_B = 1</math></p>	

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<b>SBF</b>	
<ul style="list-style-type: none"> <li>• <math>B</math> vector is replaced by a vector of counters <math>C_1, C_2, \dots, C_m</math> <ul style="list-style-type: none"> <li>• <math>C_i</math> is the sum of <math>f_x</math> values for elements <math>x \in S</math> mapping to <math>i</math></li> </ul> </li> <li>• Approximations of <math>f_x</math> are stored into                             <math display="block">C_{h_1(x)}, C_{h_2(x)}, \dots, C_{h_k(x)}</math> </li> <li>• Due to conflicts, the <math>C_i</math> provide approximations...</li> </ul> <div style="text-align: right; margin-top: 10px;"> <span style="background-color: yellow; border-radius: 50%; padding: 5px 15px; font-weight: bold;">Upper bounds</span> </div>	

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<b>The Minimum Selection</b>	
<div style="text-align: center; margin-bottom: 10px;"> <math>h_{\dots}(y) = h_{\dots}(x) = i - 1</math> </div> <ul style="list-style-type: none"> <li>• <math>C_{i-1}</math> is not a good approximation of <math>f_x</math> (neither of <math>f_y</math>)</li> <li>• <math>C_i</math> is an exact approximation of <math>f_x</math></li> <li>• <math>C_{j+1}</math> is an exact approximation of <math>f_z</math></li> </ul>	

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## Insertion and Deletion

- insertion is simple
  - increase each counter by 1
  - ...
  - for each  $h$  in  $H$  do  
     $C[h(x)] = C[h(x)] + 1$ ;
  - done
  - ...
- deletion is simple
  - decrease each counter by 1
- search for an element  $x$ 
  - return the *Minimum Selection* (MS) value  
 $m_x = \min\{C_{h_1(x)}, C_{h_2(x)}, \dots, C_{h_k(x)}\}$

Upper bounds

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Recurring minimum for  
improving the estimate  
+ 2 SBF

## On the error of SBF

- The error is the same as for Bloom Filters

**Theorem**  
For all  $x$ , it is  $f_x \leq m_x$ . Furthermore  $f_x \neq m_x$  with probability  
$$E_{SBF} = \varepsilon \approx (1 - p)^k$$

**Proof.**  
The case  $m_x < f_x$  cannot happen.  
The event  $m_x > f_x$  is "all counters  $C_{h_i(x)}$  have a collision", that corresponds to a "false positive" event of classical BF.  $\square$

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<h2>Pattern Matching</h2>	
<p>A set of objects whose keys are complex and time-costly to be compared (e.g. URLs, matrices, MP3,...).</p> <ul style="list-style-type: none"><li>• Use BF to reduce the number of explicit comparisons.</li><li>• Effective in hierarchical memories.</li><li>• Example on Dictionary matching [Bloom '70].</li></ul>	

## Set Intersection

We have two machines  $M_A$  and  $M_B$  each storing a set of items  $A$  and  $B$ , respectively. We wish to compute  $A \cap B$  exchanging a *small* number of bits.

Typical applications: *data replication check, distributed search engines.*

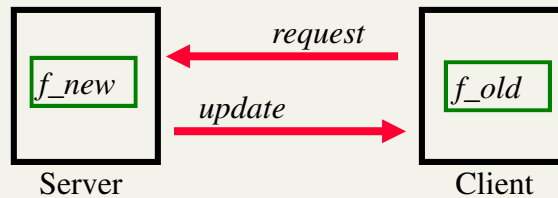
- $M_A$  sends  $BF(A)$  to  $B$ , using  $m = m_{opt} = k|A|/\ln 2$  bits.
- $M_B$  checks  $B$  into  $BF(A)$  in  $O(k|B|)$  time, and sends back *explicitly* the found items, say  $Q$ . Note that  $Q \supseteq A \cap B$ .
- $M_A$  computes  $Q \cap A$ , and returns it.

The bit-cost is  $\frac{k|A|}{\ln 2} + (|A \cap B| + |B|0.5^k) \log |U| \ll |A| \log |U|$ .  
Good for long keys,

## Web Algorithmics

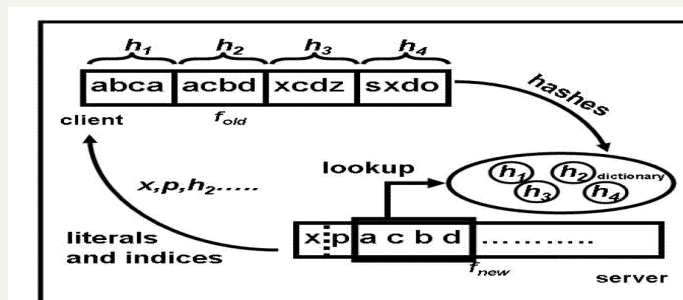
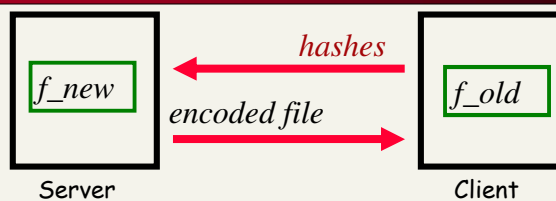
### File Synchronization

## File sync: The problem

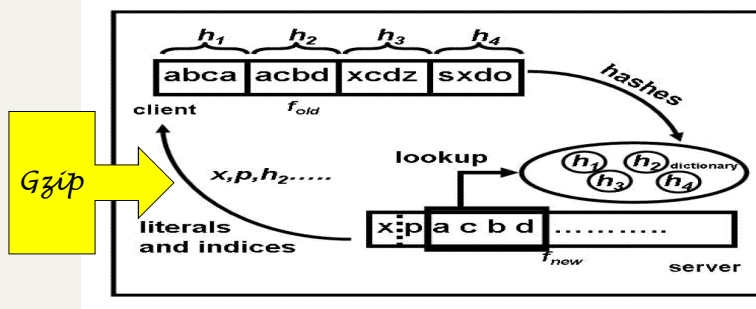


- **client** wants to update an out-dated file
- server has new file but does not know the old file
- update without sending entire  $f_{new}$  (using *similarity*)
- *rsync*: file sync tool, distributed with Linux

## The rsync algorithm



## The rsync algorithm (contd)



- simple, widely used, **single** roundtrip
- optimizations: 4-byte **rolling hash** + 2-byte **MD5**, *gzip* for literals
- choice of block size problematic (*default*:  $\max\{700, \sqrt{n}\}$  bytes)
- not good in theory: **granularity of changes may disrupt use of blocks**