

# **Big Data Analytics, The Class**

<b>Goal:</b> General A <i>model</i> or <i>summariza</i>	izations <i>tion</i> of the data.
Data Frameworks	Algorithms and Analyses
Hadoop File System Spark Streaming MapReduce Tensorflow	Similarity Search Hypothesis Testing Link Analysis Recommendation Systems Deep Learning

### **Finding Similar Items**



- There are many applications where we desire finding similar items to a given example.
- For example:
  - Document Similarity:
    - Mirrored web-pages
    - Plagiarism; Similar News
  - Recommendations:
    - Online purchases
    - Movie ratings
  - Entity Resolution: matching one instance of a person with another
  - Fingerprint Matching: finding the most likely matches in a larg dataset of matches.

### **Finding Similar Items: Topics**

- Shingling
- Minhashing
- Locality-sensitive hashing
- Distance Metrics

We will cover the following methods for finding similar items.

The first 3 make up a pipeline of techniques, culminating in LSH for rapidly matching items over a large search space. Similarity in these cases all comes down to a jaccard set similarity.

Distance metrics introduces a different set of common approaches to assessing similarity between items, assuming one has some features (quantities describing describing them).

### **Document Similarity**

**Challenge:** How to represent the document in a way that can be efficiently encoded and compared?

The first challenge for efficiently searching for similar items is simply how to represent an item.



If we can represent an item (a document in this case) simply as a set, a very simple representation, then we can look at overlap in sets as similarity.



A very easy way to get sets from all documents and many other file types is simply shingles. Take sequences of k characters in a row.



We would expect similar document to have similar shingles.

In practice using shingles of size 5 to 10 is more ideal to make it less likely to randomly match shingles between 2 documents.

Goal: Con	vert documents to sets
	Large enough that any given shingle appearing a document is highly unlikely (e.g. < .1% chance) Can hash large shingles to smaller (e.g. 9-shingles into 4 bytes)
	<ul> <li>Can also use words (aka n-grams).</li> <li>Similar documents have many common shingles</li> </ul>
	<ul> <li>Changing will fis or order has minimal effect.</li> <li>In practice use 5 &lt; k &lt; 10</li> </ul>

Generally, we want elements in our sets (i.e. shingles) to match with about 1 in 1000 probability.

The larger generally the better for this purpose and we can even hash shingles to reduce their size a bit.

**Problem:** Even if hashing, sets of shingles are large (e.g. 4 bytes => 4x the size of the document).

However, such a representation, even when hashed, still enlarges the document rather than reduces it and we want to be able to search over millions to billions of these quickly. If you consider a character as a byte then even hashing 9grams (9 bytes) down to 4 bytes has the potential to make a document 4x its original size.

Goal: Convert sets to shorter ids, signatures

While shingles gives us a simple way to turn a document into a set, we need a way to make that set representation smaller. This is where minhashing comes in.

### Goal: Convert sets to shorter ids, "signatures"



Let's go ahead and define how we will compute similarity based on a set:

We can use Jaccard Similarity: The amount of overlap divided by the total elements of the union.

In this way, similarity is basically a percentage of the total number of elements that are shared.

It has intuitive properties such as if one document is larger and thus has more elements in its set that will have the effect of shrinking the amount of similarity unless they other document contains many of the same elements.

We will call "characteristic matrix" the actual type of data structure we use to represent these sets. It's simply a binary matrix with sets (i.e. documents) as columns and shingles (i.e. elements) as rows.

In practice, the characteristic matrix will be very sparse -- remember we want about a 1 in 1000 chance of a particular shingle to appear.

	<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>
ab	1	1
bc	0	1
de	1	0
ah	1	1
ha	0	0
ed	1	1
са	0	1

Jaccard Similarity:

$$sim(S_1, S_2) = \frac{S_1 \cap S_2}{S_1 \cup S_2}$$

Latex equation:  $sim(S_1, S_2) = \frac{S_1 \cos S_2}{S_1 \cos S_2}$ 

Let's start to work with an example charactertistic matrix of two documents.

What would be the similarity?

#### Characteristic Matrix:

	S <sub>1</sub>	<i>S</i> <sub>2</sub>	
ab	1	1	* *
bc	0	1	*
de	1	0	*
ah	1	1	**
ha	0	0	
ed	1	1	**
са	0	1	*

Jaccard Similarity:

$$sim(S_1, S_2) = \frac{S_1 \cap S_2}{S_1 \cup S_2}$$

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One way to quick algorithm to calculate is simply to sum the rows.



Jaccard Similarity:

$$sim(S_1, S_2) = \frac{S_1 \cap S_2}{S_1 \cup S_2}$$

 $sim(S_1, S_2) = 3 / 6$ # both have / # at least one has

and divide the number of 2s by the number of 1s. (i.e. 3/6 in this case)

Notice we only care about when one of them is 1.

Problem: Even if hashing shingle contents, sets of shingles are large
e.g. 4 byte integer per shingle: assume all unique shingles,
=> 4x the size of the document
(since there are as many shingles as characters and 1byte per char).

So, keeping Jaccard Similarity in mind, how do we get this characteristic matrix smaller?

Miı	nh	a	sł	nir	ng	Goal: Convert sets to shorter ids, "signatures"
Chara	acte	eris	stic	Ma	atrix	c: <i>X</i>
		<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>	
	ab	1	0	1	0	
	bc	1	0	0	1	
	de	0	1	0	1	
	ah	0	1	0	1	
	ha	0	1	0	1	
	ed	1	0	1	0	
	ca	1	0	1	0	
(Leskove	ec at a	l., 201	l4; htt	p://ww	/w.mm	ds.org/)

We want to create a shorter id a "signature" from the larger characteristic matrix

Mi	Minhashing						Goal: Convert sets to shorter ids, "signatures"		
Chara	Characteristic Matrix: X				atri	<b>x</b> : X	<b>Approximate Approach:</b> 1) Instead of keeping whole characteristic matrix, just keep first row where 1 is encountered.		
		<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>	-	2) Shuffle and repeat to get a "signature" for each set.		
	ab	1	0	1	0				
	bc	1	0	0	1				
	de	0	1	0	1				
	ah	0	1	0	1				
	ha	0	1	0	1				
	ed	1	0	1	0				
	ca	1	0	1	0	1			
(Leskove	ha ed ca	0 1 1 ., 20'	1 0 0	0 1 1 p://wv	1 0 0 vw.mm	nds.org/	)		

Well let's take an extreme approach. What if we only represented the Set by a single integer?

We could just keep the row number where the first element was non-zero.

Min	h	a	sh	nir	ng		Goal: Convert sets to shorter ids, "signatures"
Charac	cte	ris 1	tic 3	Ma 1	atrix 2	<b>x</b> : X	<u>Approximate Approach:</u> 1) Instead of keeping whole characteristic matrix, just keep first row where 1 is encountered.
		<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>		2) Shuffle and repeat to get a "signature" for each set.
a	ıb	1	0	1	0	-	
b	ос	1	0	0	1		
d	le	0	1	0	1		
a	ıh	0	1	0	1		
h	na	0	1	0	1		
e	ed	1	0	1	0		
с	a	1	0	1	0		
(Leskovec a	at al.	, 201	4; htt	p://wv	/w.mm	ids.org/)	

Here is what we would get: set 1 and set 3 would actually get the same integer, while 2 and 4 would each have a different.

Well set 1 and set 3 do happen to be quite similar: Their Sim is 3/4

In fact, if you think about it, given a random ordering of the rows, what is the probability that both of their first non-zero row happens to be the same? <sup>3</sup>/<sub>4</sub> in 3 of the 4 possible rows that have at least a 1 (ab, bv, ed, and ca) only 1 of them being first wouldn't be a match (bc).

Mir	Minhashing								onve	ert so	ets t	o shorter ids, "signatures"
Chara	acte	ris 1 $S_1$	$S_2$	M: 1 <i>S</i> <sub>3</sub>	$\frac{1}{S_4}$	<b>x</b> : <i>X</i>	<u>Apr</u> 1) lr kee 2) S	prox nste p fir	ad o st ro	of ke by w and	Appi epir /her/ <b>rep</b>	roach: ng whole characteristic matrix, just e 1 is encountered. eat to get a "signature".
	ab	1	0	1	0			$\bigcirc$				
	bc	1	0	0	1			$\langle \mathbf{Z} \rangle$				
	de	0	1	0	1	-		S <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>	
	ah	0	1	0	1	-	ah	0	1	0	1	
	ha	0	1	0	1	-	ca	1	0	1	0	
	ed	1	0	1	0	-	ed	1	0	1	0	
	ca	1	0	1	0	-	de	0	1	0	1	•••
							ab	1	0	1	0	
(Leskove	ec at a	., 201	14; htt	p://wv	vw.mn	nds.org/)	bc	1	0	0	1	

In reality of course, a single integer is not going to be enough but we can repeat this a few times. Here's an example after we shuffle.

Now both pairs S1 - S3 AND S2 S4 match. S2 and S4 also have a sim of  $\frac{3}{4}$ . If we just asked at this point how much did these 2-integer signatures match, we'd find 100% for S1-S3 and 50% for S2-S4... one overestimates; one underestimates... This can continue in order to make a more and more accurate signature that matches with the same probability as the Jaccard Similarity.



Here is what the signatures look like so far.

We're going to try to produce a "signature matrix" as the output of minhashing, where each column is a signature.



One downside of how we've discuss this is the time it would take to keep reshuffling rows, but there's really no need to do that.

Shuffle is just the conceptual way to think about this when in fact we can use hash functions to give us a random order of rows to look at.

#### Characteristic Matrix:

	<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>
ab	1	0	1	0
bc	1	0	0	1
de	0	1	0	1
ah	0	1	0	1
ha	0	1	0	1
ed	1	0	1	0
ca	1	0	1	0

### Minhash function: h

• Based on permutation of rows in the characteristic matrix, *h* maps sets to first row where set appears.

(Leskovec at al., 2014; http://www.mmds.org/)









 $sim(S_1, S_2) = \frac{S_1 \log S_2}{S_1 \log S_2}$ 



 $sim(S_1, S_2) = \frac{S_1 \log S_2}{S_1 \log S_2}$ 

#### Characteristic Matrix:

		<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	S <sub>3</sub>	<i>S</i> <sub>4</sub>
3	ab	1	0	1	0
4	bc	1	0	0	1
7	de	0	1	0	1
6	ah	0	1	0	1
1	ha	0	1	0	1
2	ed	1	0	1	0
5	са	1	0	1	0

Minhash function: h

Based on permutation of rows in the • characteristic matrix, h maps sets to rows.

### Signature matrix: M

Record first row where each set • had a 1 in the given permutation

	$S_1$	$S_2$	$S_{3}$	$S_4$
$h_1$	2	1	2	1

- $h_1(S_1) = ed$  #permuted row 2  $h_1(S_2) = ha$  #permuted row 1  $h_1(S_3) = ed$  #permuted row 2  $h_1(S_4) = ha$  #permuted row 1

 $sim(S_1, S_2) = \frac{S_1 \cos S_2}{S_1 \cos S_2}$ 





#### Characteristic Matrix:

			<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	S <sub>3</sub>	<i>S</i> <sub>4</sub>
4	3	ab	1	0	1	0
2	4	bc	1	0	0	1
1	7	de	0	1	0	1
3	6	ah	0	1	0	1
6	1	ha	0	1	0	1
7	2	ed	1	0	1	0
5	5	са	1	0	1	0

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$h_1$	2	1	2	1
$h_2$				

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#### Characteristic Matrix:

			<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	S <sub>3</sub>	<i>S</i> <sub>4</sub>
4	3	ab	1	0	1	0
2	4	bc	1	0	0	1
1	7	de	0	1	0	1
3	6	ah	0	1	0	1
6	1	ha	0	1	0	1
7	2	ed	1	0	1	0
5	5	са	1	0	1	0

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$h_1$	2	1	2	1
h <sub>2</sub>	2	1	4	1

#### $sim(S_1, S_2) = \frac{S_1 \cos S_2}{S_1 \cos S_2}$

#### Characteristic Matrix:

				$S_1$	$S_2$	S <sub>3</sub>	$S_4$
1	4	3	ab	1	0	1	0
3	2	4	bc	1	0	0	1
7	1	7	de	0	1	0	1
6	3	6	ah	0	1	0	1
2	6	1	ha	0	1	0	1
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$h_1$	2	1	2	1
$h_2$	2	1	4	1
h <sub>3</sub>				

(Leskovec at al., 2014; http://www.mmds.org/)

#### Characteristic Matrix:

				<i>S</i> <sub>1</sub>	$S_2$	<i>S</i> <sub>3</sub>	$S_4$
1	4	3	ab	1	0	1	0
3	2	4	bc	1	0	0	1
7	1	7	de	0	1	0	1
6	3	6	ah	0	1	0	1
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_	-						

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$h_1$	2	1	2	1
$h_2$	2	1	4	1
h <sub>3</sub>	1	2	1	2

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3	2	4	bc	1	0	0	1
7	1	7	de	0	1	0	1
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$h_1$	2	1	2	1
h <sub>2</sub>	2	1	4	1
h <sub>3</sub>	1	2	1	2

(Leskovec at al., 2014; http://www.mmds.org/)


 $sim(S_1, S_2) = \frac{S_1 \exp S_2}{S_1 \exp S_2}$ 

Minhashing							<b>Property of signature matrix:</b> The probability for any $h_i$ (i.e. any row), that							
Characteristic Matrix:						Thus	$h_i(S_1) = h_i(S_2)$ is the same as Sim $(S_1, S_2)$ Thus, similarity of signatures $S_1, S_2$ is the fraction of							
$S_1 S_2 S_3 S_4$						min	minhash functions (i.e. rows) in which they agree.							
1	4	3	ab	1	0	1	0			lno	9			
3	2	4	bc	1	0	0	1			S	S	S	S	
7	1	7	de	0	1	0	1	-	h	2 2	1	2 <sub>3</sub>	1 <sup>3</sup>	
6	3	6	ah	0	1	0	1		<sup>n</sup> 1	2	1	2	1	
2	6	1	ha	0	1	0	1		<i>n</i> <sub>2</sub>	2	1	4	1	
5	7	2	ed	1	0	1	0		h <sub>3</sub>	1	2	1	2	
4	5	5	са	1	0	1	0							
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 $sim(S_1, S_2) = \frac{S_1 \cos S_2}{S_1 \cos S_2}$ 



#### Error Bound?

Expect error:  $O(1/\sqrt{k})$  (k hashes) Why? Each row is a random observation of 1 or 0 (match or not) with P(match=1) = Sim(S1, S2). N = k observations

Standard deviation(*std*)? < 1 (worst case is 0.5)

	<i>S</i> <sub>1</sub>	<i>S</i> <sub>2</sub>	<i>S</i> <sub>3</sub>	<i>S</i> <sub>4</sub>
h <sub>1</sub>	2	1	2	1
h <sub>2</sub>	2	1	4	1
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Estimated Sim(S<sub>1</sub>, S<sub>3</sub>) = agree / all = 2/3

Real Sim(S<sub>1</sub>, S<sub>3</sub>) = Type a / (a + b + c) = 3/4

Try Sim(S $_{2},$  S $_{4}) and Sim(S<math display="inline">_{1},$  S $_{2})$ 

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 $sim(S_1, S_2) = \frac{S_1 \cos S_2}{S_1 \cos S_2}$ 



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Standard Error of Mean =  $std/\sqrt{N}$ 

	<i>S</i> <sub>1</sub>	$S_2$	S <sub>3</sub>	<i>S</i> <sub>4</sub>
$h_1$	2	1	2	1
h <sub>2</sub>	2	1	4	1
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In Practice

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- Can't reasonably do permutations (huge space)
- Can't randomly grab rows according to an order (random disk seeks = slow!)

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Solution: Use "random" hash functions.

- Setup:
  - Pick ~100 hash functions, hashes
  - Store M[i][s] = a potential minimum h<sub>i</sub>(r)
     *#initialized to infinity (num hashs x num sets)*

```
Solution: Use "random" hash functions.

Setup:

hashes = [getHfunc(i) for i in rand(1, num=100)]

#100 hash functions, seeded random

for i in hashes: for s in sets:

M[i][s] = np.inf #represents a potential minimum h_i(r); initially infinity

Algorithm ("efficient minhashing"):

for r in rows of cm: #cm is characteristic matrix

compute h_i(r) for all i in hashes #precompute 100 values

for each set s in sets:

if cm[r][s] == 1:

for i in hashes: #check which hash produces smallest value

if h_i(r) < M[i][s]: M[i][s] = h_i(r)
```

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Come up with example?

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**New Problem:** Even if the size of signatures are small, it can be computationally expensive to find similar pairs.

E.g. 1m documents; 1,000,000 choose 2 = 500,000,000,000 pairs!

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E.g. 1m documents; 1,000,000 choose 2 = 500,000,000,000 pairs!

(1m documents isn't even "big data")

Come up with example?



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If we wanted the similarity for all pairs of documents, could anything be done?

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Approach: Hash multiple times over subsets of data: similar items are likely in the same bucket once.

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Approach from MinHash: Hash columns of signature matrix

→ Candidate pairs end up in the same bucket.

















## Probabilities of agreement, Example

- 100,000 documents
- 100 random permutations/hash functions/rows
   => if 4byte integers then 40Mb to hold signature matrix
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- Want 80% Jaccard Similarity ; for any row  $p(S_1 == S_2) = .8$

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   => still 100k choose 2 is a lot (~5billion)
- 20 bands of 5 rows
- Want 80% Jaccard Similarity ; for any row  $p(S_1 == S_2) = .8$

 $P(S_1 = S_2 | b^{(5)})$ : probability S1 and S2 agree within a given band










Pipeline gives us a way to find *near-neighbors* in *high-dimensional space* based on Jaccard Distance (1 - Jaccard Sim).



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Typical properties of a distance metric, *d*:

d(a, a) = 0

d(a, b) = d(b, a)

 $d(a, b) \le d(a,c) + d(c,b)$ 



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There are other metrics of similarity. e.g:

- Euclidean Distance
- Cosine Distance

. . .

- Edit Distance
- Hamming Distance

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There are other metrics of similarity. e.g:

• Euclidean Distance

distance
$$(X, Y) = \sqrt{\sum_{i=1}^{n} (x_i - y_i)^2}$$
 ("L2 Norm")

Cosine Distance

. . .

- Edit Distance
- Hamming Distance

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# Locality Sensitive Hashing - Theory

LSH Can be generalized to many distance metrics by converting output to a probability and providing a lower bound on probability of being similar.

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LSH Can be generalized to many distance metrics by converting output to a probability and providing a lower bound on probability of being similar.

E.g. for euclidean distance:

- Choose random lines (analogous to hash functions in minhashing)
- Project the two points onto each line; match if two points within an interval

#### Side Note on Generating Hash Functions:

#### What hash functions to use?

Start with 2 decent hash functions

e.g.  $h_a(x) = ascii(string) \% large_prime_number$  $h_b(x) = (3*ascii(string) + 16) \% large_prime_number$ 

Add together multiplying the second times i:

 $\begin{aligned} h_i(x) &= h_a(x) + i^* h_b(x) \ \% \ |BUCKETS| \\ e.g. \ h_5(x) &= h_a(x) + 5^* h_b(x) \ \% \ 100 \end{aligned}$ 

https://www.eecs.harvard.edu/~michaelm/postscripts/rsa2008.pdf

Popular choices: md5 (fast, predistable); mmh3 (easy to seed; fast)