

Data Mining and Analysis

Finding similar items

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Texas A&M University

Department of Computer Science & Engineering

Prof. James Caverlee

Resources

MMDS Chapter 3 + slides

[http://i.stanford.edu/~ullman/mmds/
ch3n.pdf](http://i.stanford.edu/~ullman/mmds/ch3n.pdf)

[http://www.mmds.org/mmds/v2.1/ch03-
lsh.pdf](http://www.mmds.org/mmds/v2.1/ch03-lsh.pdf)

Carlos Castillo course on Data Mining [[https://
github.com/chatox/data-mining-course](https://github.com/chatox/data-mining-course)]

Example

- **Assume the following case:**
 - Suppose 100,000 columns of M (100k docs)
 - Signatures of 100 integers (rows)
(Therefore, signatures take 40Mb)
 - Choose $b = 20$ bands of $r = 5$ integers/band
- **Goal:** Find pairs of documents that are at least $s = 0.8$ similar

Example: Suppose $\text{sim}(C_1, C_2) = 0.8$

- Find pairs of $\geq s=0.8$ similarity, set $b=20$, $r=5$
- Since $\text{sim}(C_1, C_2) \geq s$, we want C_1, C_2 to be a candidate pair:
 - We want them to hash to at least 1 common bucket (at least one band is identical)
- Probability C_1, C_2 identical in one particular band: $(0.8)^5 = 0.328$
- Probability C_1, C_2 are **not** similar in all of the 20 bands: $(1-0.328)^{20} = 0.00035$
 - i.e., about 1/3000th of the 80%-similar column pairs are **false negatives** (we miss them)
 - We would find **99.965%** pairs of truly similar documents

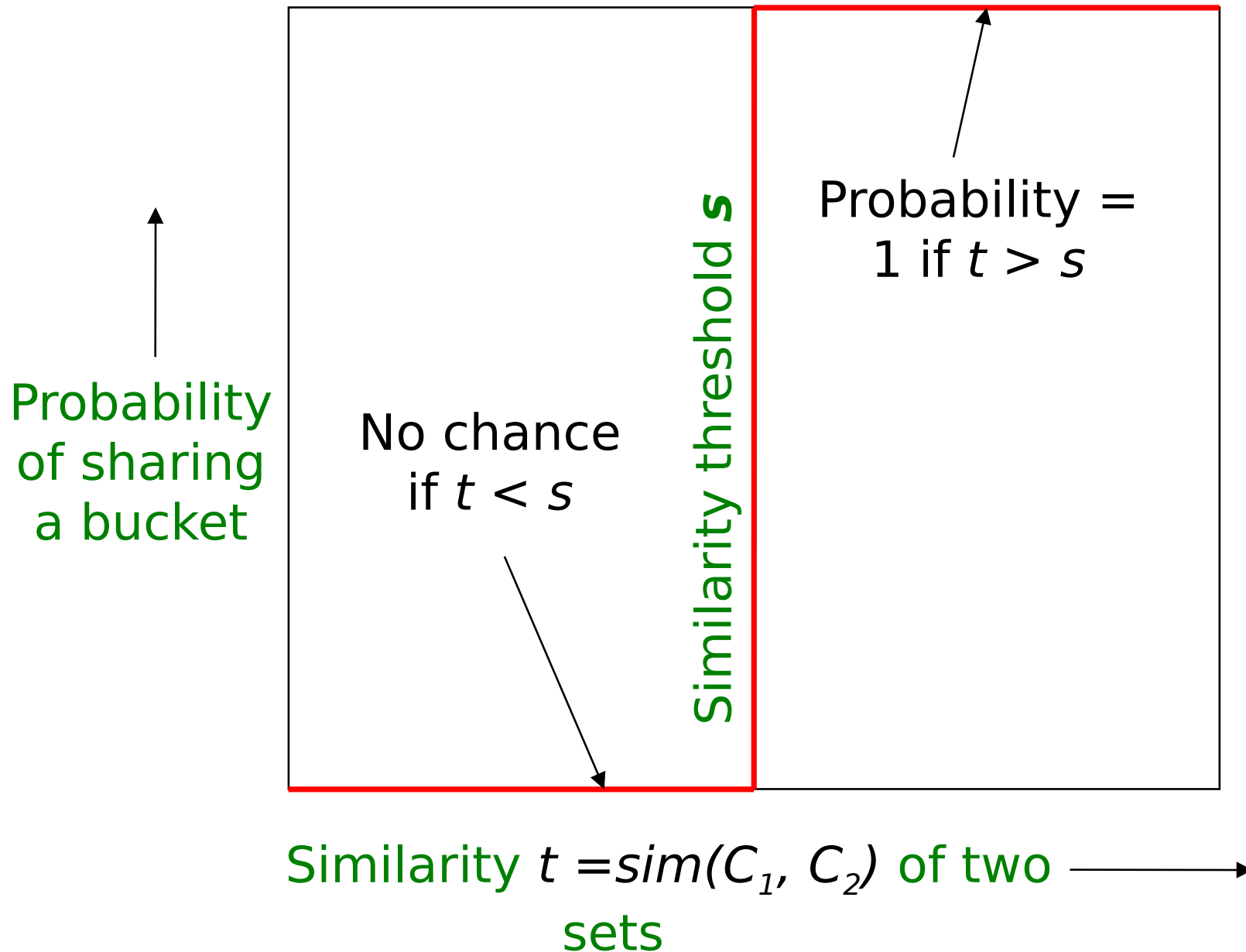
Example: Suppose $\text{sim}(C_1, C_2) = 0.3$

- Find pairs of $\geq s=0.8$ similarity, set $b=20$, $r=5$
- Since $\text{sim}(C_1, C_2) < s$ we want C_1, C_2 to hash to NO common buckets (all bands should be different)
- Probability C_1, C_2 identical in one particular band:
 $(0.3)^5 = 0.00243$
- Probability C_1, C_2 identical in at least 1 of 20 bands: $1 - (1 - 0.00243)^{20} = 0.0474$
- In other words, approximately 4.74% pairs of docs with similarity 0.3% end up becoming **candidate pairs**
- They are **false positives** since we will have to examine them (they are candidate pairs) but then it will turn out their similarity is below threshold s

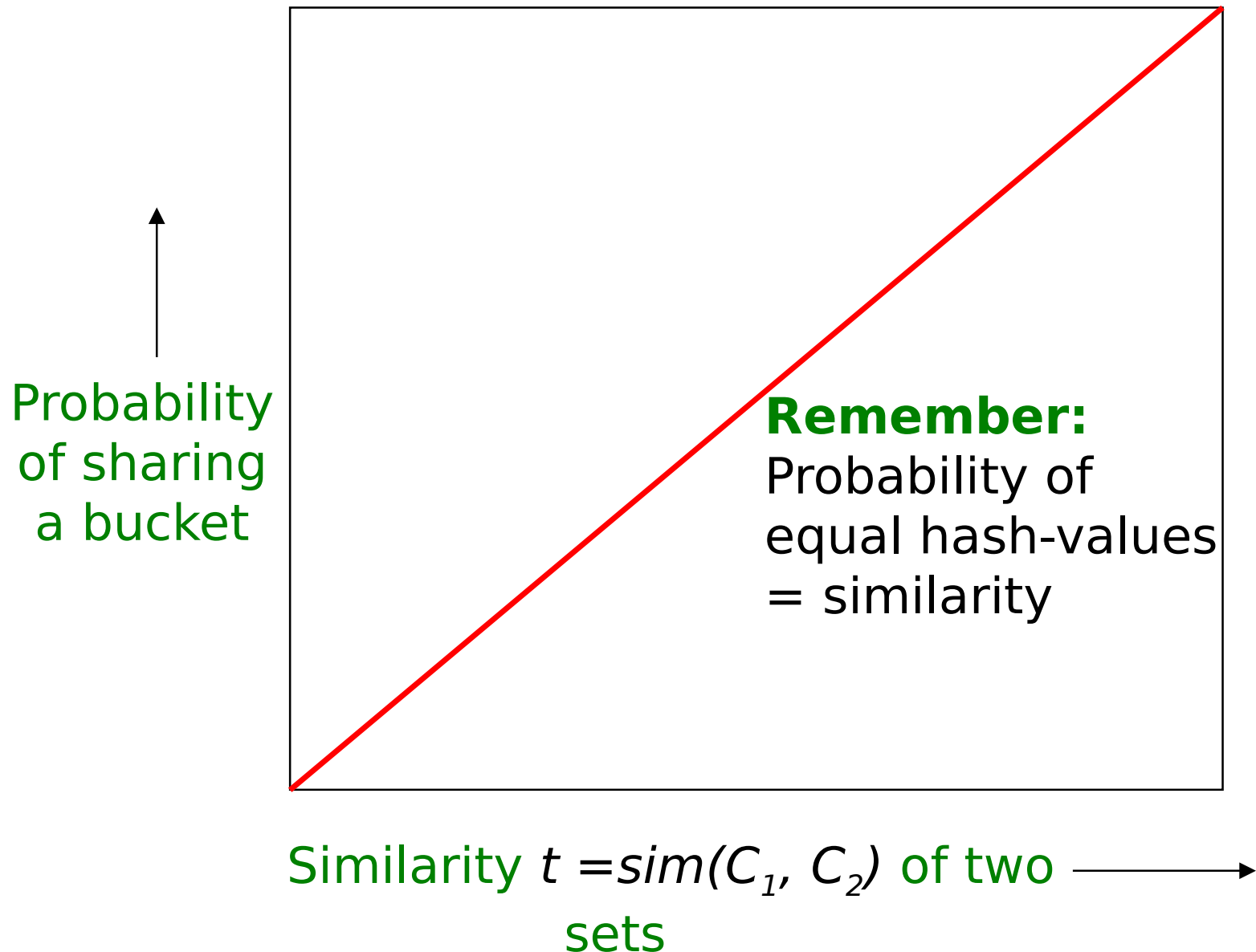
LSH involves a trade-off

- **Pick:**
 - The number of Min-Hashes (rows of M)
 - The number of bands b , and
 - The number of rows r per band to balance false positives/negatives
- **Example:** If we had only 15 bands of 5 rows, the number of false positives would go down, but the number of false negatives would go up

LSH: what we want



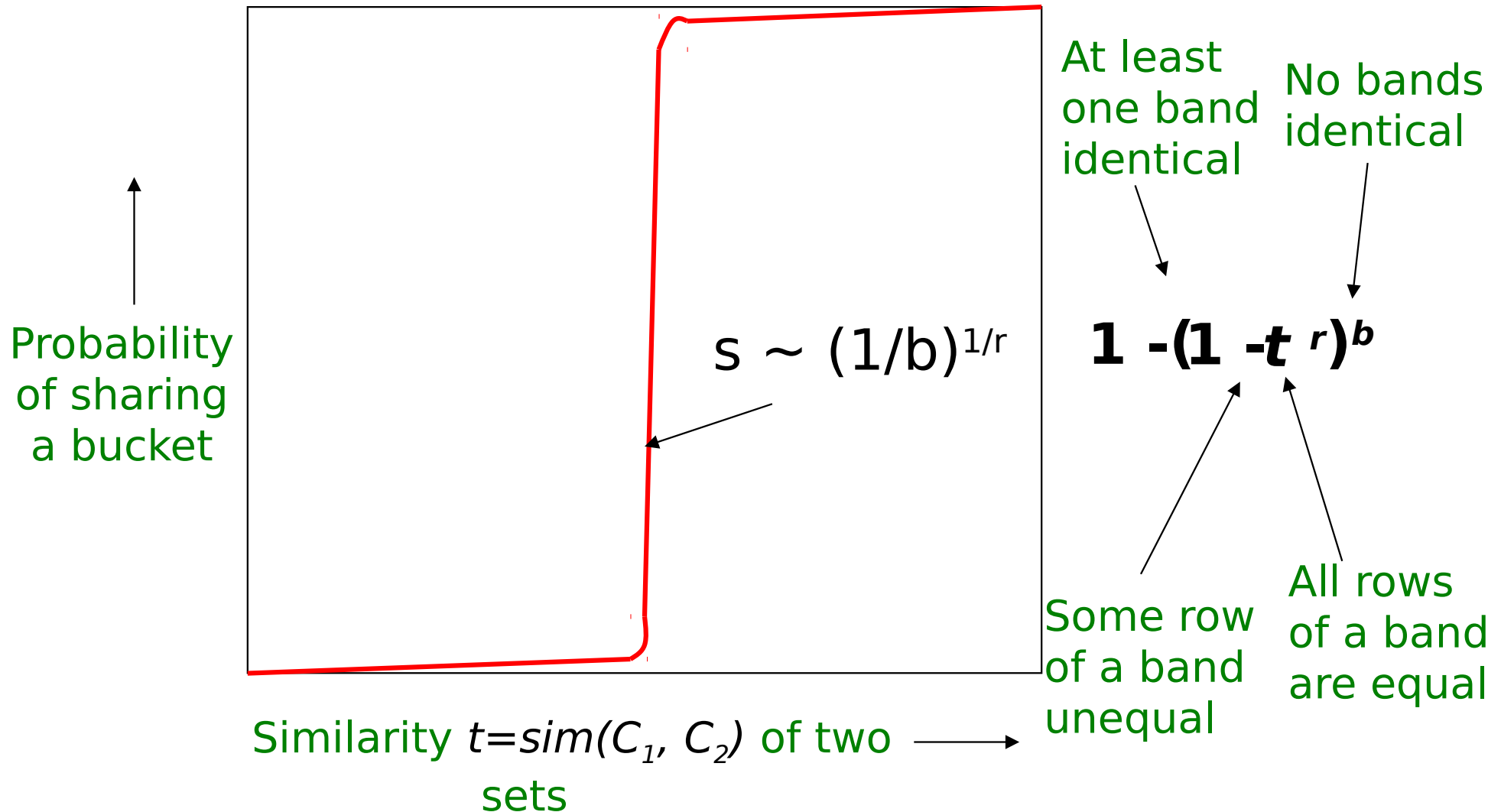
What 1 band of 1 row gives you



b bands, r rows/band

- Columns C_1 and C_2 have similarity t
- Pick any band (r rows)
 - Prob. that all rows in band equal =
 - t^r
 - Prob. that some row in band unequal =
 - $1 - t^r$
- Prob. that no band identical =
 - $(1 - t^r)^b$
- Prob. that at least 1 band identical =
 - $1 - (1 - t^r)^b$

What b bands of r rows gives you



Example: $b=20$, $r=5$

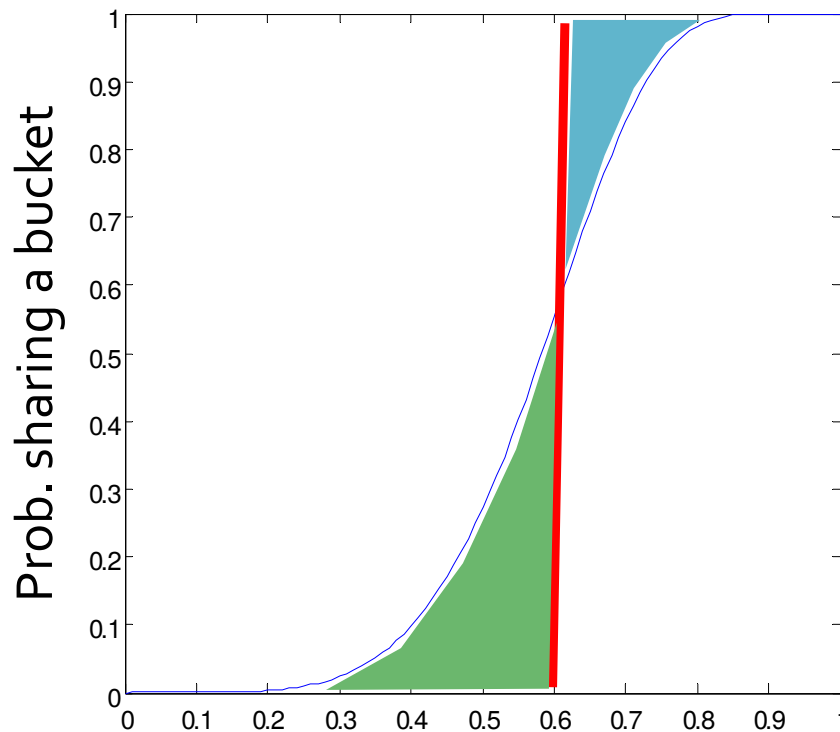
Similarity threshold s

Probability that at least
1 band is identical:

s	$1-(1-s^r)^b$
.2	.006
.3	.047
.4	.186
.5	.470
.6	.802
.7	.975
.8	.9996

Picking r and b : the S curve

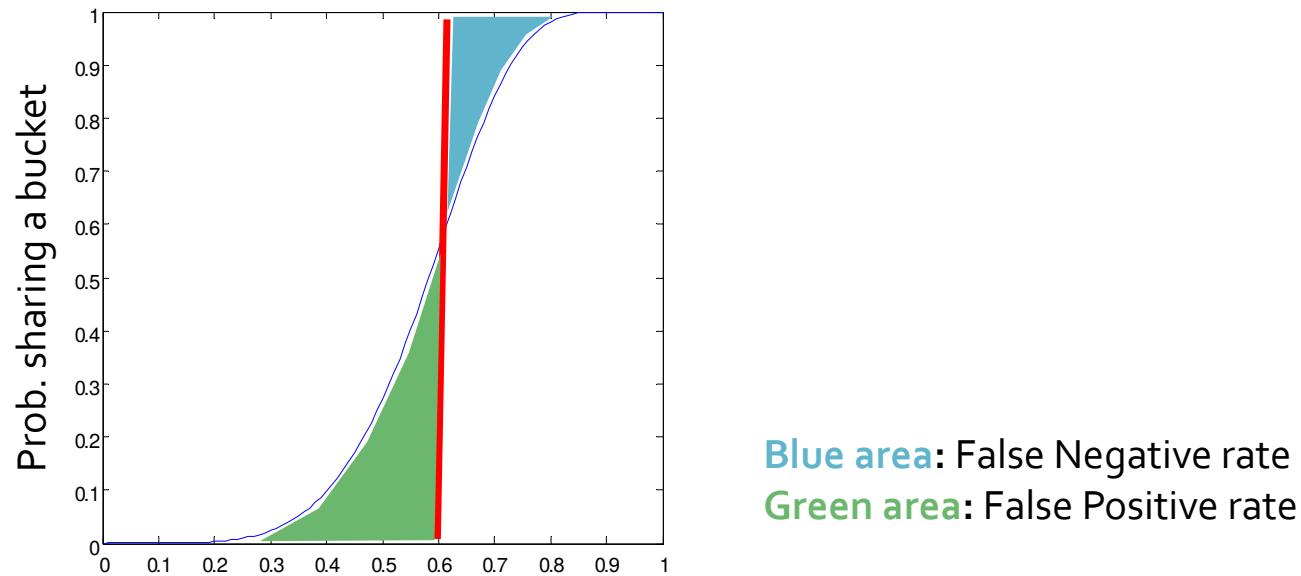
Picking r and b to get the best S-curve
50 hash-functions ($r=5$, $b=10$)



Blue area: False Negative rate
Green area: False Positive rate

Picking r and b to get the best S-curve

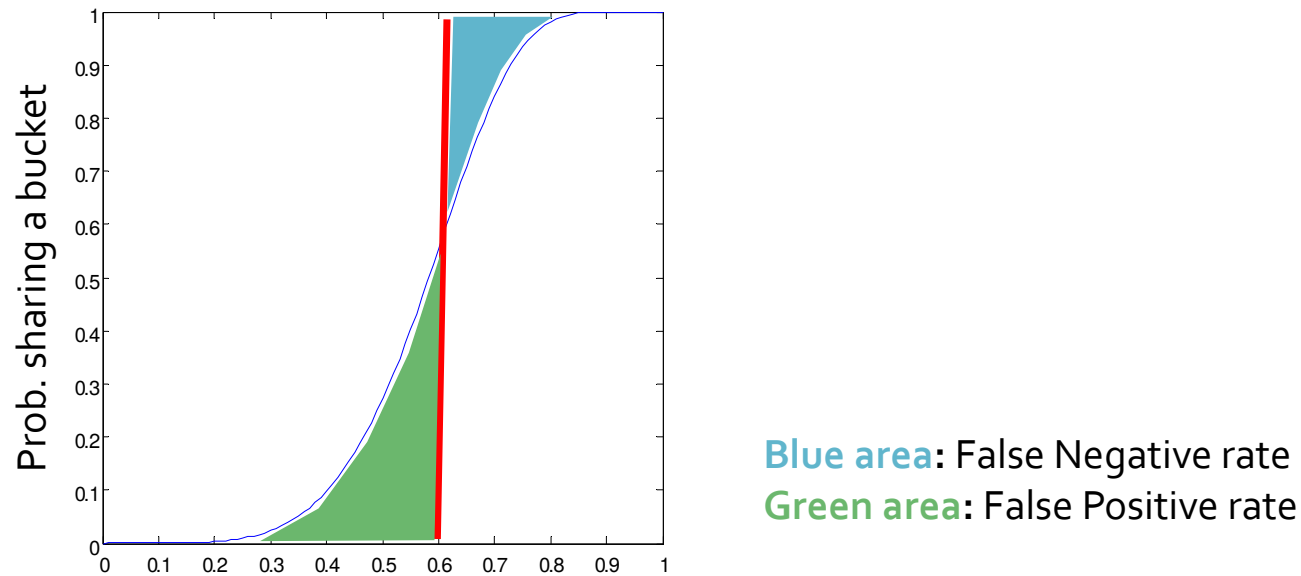
50 hash-functions ($r=5$, $b=10$)



Blue area X: False Negative rate These are pairs with $\text{sim} > s$ but the X fraction won't share a band and then will never become candidates. This means we will never consider these pairs for (slow/exact) similarity calculation!

Picking r and b to get the best S-curve

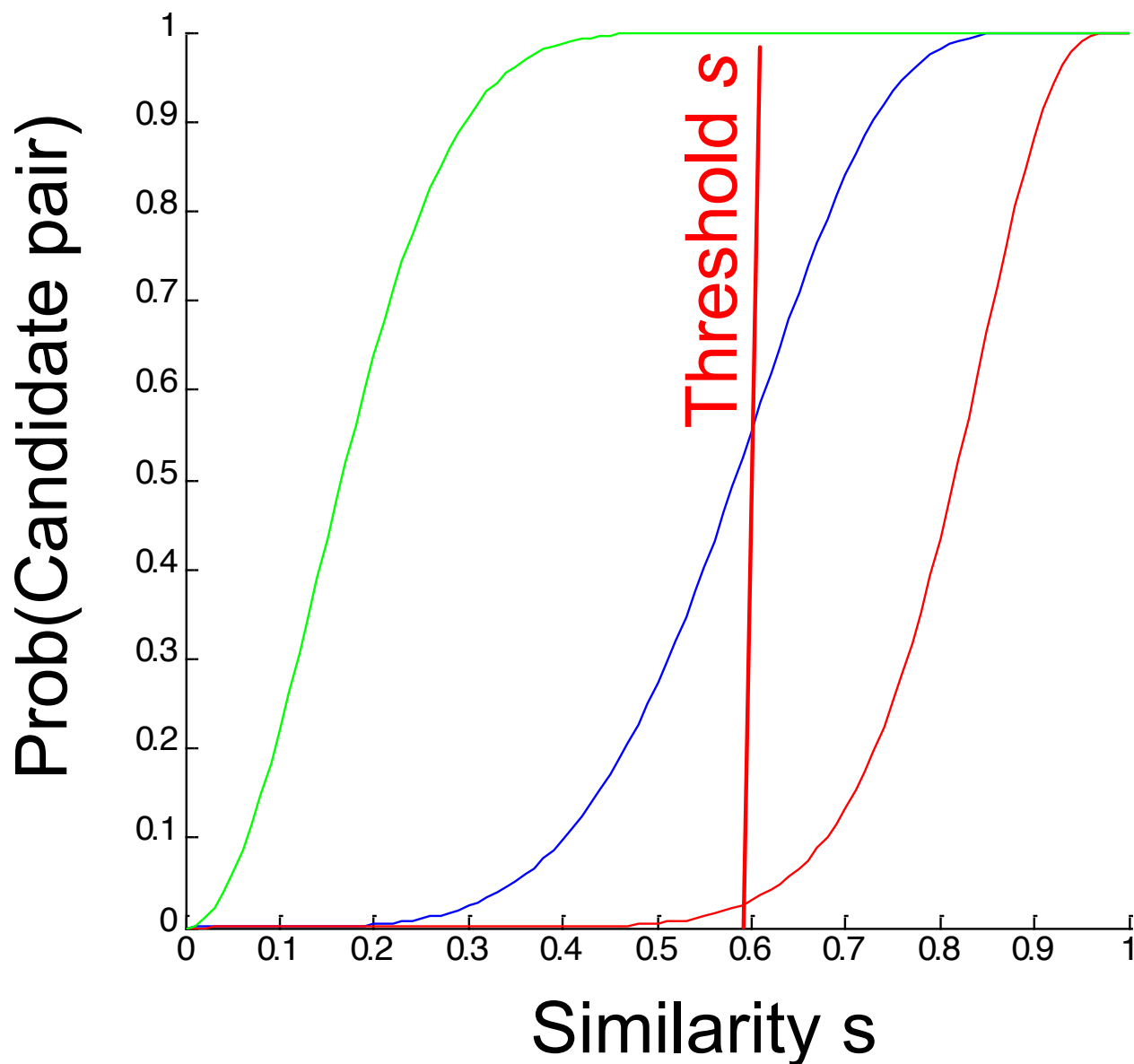
50 hash-functions ($r=5$, $b=10$)



Green area Y: False positive rate.

These are pairs with $\text{sim} < s$ but we will consider them as candidates. This is not too bad, we will consider them for (slow/exact) similarity computation and discard them

Suppose we have 50 hash functions ($r * b = 50$)



$r=2, b=25$

$r=5, b=10$

$r=10, b=5$

LSH Summary

Tune M , b , r to get almost all pairs with similar signatures, but eliminate most pairs that do not have similar signatures

Check in main memory that candidate pairs really do have similar signatures

Optional: In another pass through data, check that the remaining candidate pairs really represent similar documents