## **String Similarity Join**

Dong Deng

#### Real World Data is Dirty

• Misspellings of the query "britney spears" on Google

```
488941 britney spears
                            29 britent spears
                                                     9 brinttany spears
40134 brittany spears
                            29 brittnany spears
                                                     9 britanay spears
36315 brittney spears
                           29 britttany spears
                                                     9 britinany spears
24342 britany spears
                            29 btiney spears
                                                     9 britn spears
 7331 britny spears
                            26 birttney spears
                                                     9 britnew spears
                                                     9 britneyn spear G
 6633 briteny spears
                            26 breitney spears
                                                     9 britrney spears
 2696 britteny spears
                           26 brinity spears
                           26 britenay spears
 1807 briney spears
                                                     9 brtiny spears
                           26 britneyt spears
 1635 brittny spears
                                                     9 brtittney spears
 1479 brintey spears
                            26 brittan spears
                                                     9 brtny spears
 1479 britanny spears
                            26 brittne spears
                                                     9 brytny spears
                                                     9 rbitney spears
 1338 britiny spears
                            26 btittany spears
 1211 britnet spears
                            24 beitney spears
                                                     8 birtiny spears
 1096 britiney spears
                            24 birteny spears
                                                     8 bithney spears
  991 britaney spears
                            24 brightney spears
                                                     8 brattany spears
  991 britnay spears
                            24 brintiny spears
                                                     8 breitny spears
  811 brithney spears
                            24 britanty spears
                                                     8 breteny spears
  811 brtiney spears
                            24 britenny spears
                                                     8 brightny spears
  664 birtney spears
                            24 britini spears
                                                     8 brintay spears
                            24 britnwy spears
  664 brintney spears
                                                     8 brinttey spears
                            24 brittni spears
  664 briteney spears
                                                     8 briotney spears
```

http://marc.merlins.org/linux/talks/google/britney.html

### **Set Similarity Functions**

Overlap Size

$$overlap(x,y) = |x \cap y|$$

Jaccard Similarity

$$J(x,y) = \frac{|x \cap y|}{|x \cup y|}$$

$$x = \{A,B,C,D,E\}$$
  
 $y = \{B,C,D,E,F\}$ 

$$overlap(x, y) = 4$$

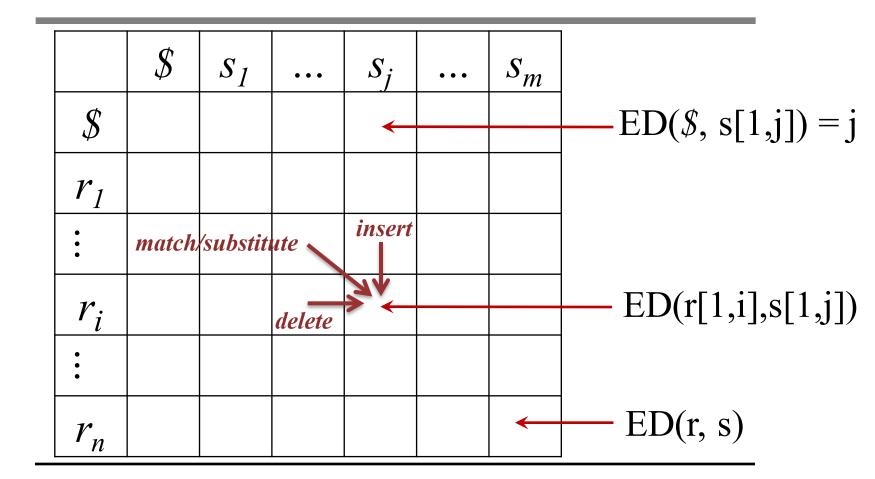
$$J(x, y) = 4/6 = 0.67$$

#### **String Similarity Function**

- Edit Distance ED(r, s): the minimum number of edit operations (insertion/deletion/substitution) needed to transform r to s.
- For example: ED(hilton, huston) = 2

• Edit Similarity: EDS(r, s) =  $1 - \frac{ED(r,s)}{\max(|r|,|s|)}$ 

```
Calculating ED(r,s)
Let r_n and s_m be the last characters in r and s.
 a) Match r_n and s_m
        ED(r,s) = 0 + ED(r[1,n-1],s[1,m-1])
 b) Substitute r<sub>n</sub> with s<sub>m</sub>
        ED(r,s) = 1 + ED(r[1,n-1],s[1,m-1])
 c) Delete r<sub>n</sub>
        ED(r,s) = 1 + ED(r[1,n-1],s[1,m])
 d) Insert s<sub>m</sub>
        ED(r,s) = 1 + ED(r[1,n],s[1,m-1])
```



	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1							
t	2							
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	012	2	3	4	5	6	7
b	12	0						
t	2							
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0-	<b>≯</b> 1					
t	2							
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0-	<b>≯</b> 1→	<b>2</b>				
t	2							
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0	1	2	3	4	5	6
t	2							
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0	1	2	3	4	5	6
t	2	1	1	2	3	4	5	6
i	3							
n	4							
e	5							
y	6							

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0	1	2	3	4	5	6
t	2	1	1	2	3	4	5	6
i	3	2	2	1	2	3	4	5
n	4	3	3	2	2	2	3	4
e	5	4	4	3	3	3	2	3
y	6	5	5	4	4	4	3	2

	<b>\$</b>	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0	1	2	3	4	5	6
t	2	1	1	2	3	4	5	6
i	3	2	2	1	2	3	4	5
n	4	3	3	2	2	2	3	4
e	5	4	4	3	3	3	2	3
y	6	5	5	4	4	4	3	2

	\$	b	r	i	t	n	e	y	From s='btiney' to r='britney'
\$	0	1	2	3	4	5	6	7	Match s[1]='b' $\rightarrow b$
b	1	0	1 subsi	2 titute	3	4	5	6	Substitute s[2]='r' with 't' $\rightarrow bt$
t	2	1	1.	2	3	4	5	6	Match $s[3] = 'i' \rightarrow bti$
i	3	2	2	1 <u>de</u>	lete >2	3	4	5	Delete s[4]='t' $\rightarrow bti$
n	4	3	3	2	2	2	3	4	Match s[5]='n' $\rightarrow$ btin
e	5	4	4	3	3	3	2	3	Match s[6]='e' $\rightarrow$ btine
y	6	5	5	4	4	4	3	2	Match s[7]='y' $\rightarrow$ btiney = r

#### What's the difference?

Functions (normalization)	Edit Distance Edit Similarity	Overlap Size Jaccard Similarity
Input	Sequences	Sets
Example Representations	DNA, String, Time Series	Image, Document, Vector, Friend List

### **String Similarity Join**

- Input:
  - A collection of strings S
  - A threshold  $\tau$
- Output:
  - All string pairs  $(s,r) \in S \times S$  such that  $ED(s,r) \leq \tau$

### **String Similarity Join**

• Give threshold  $\tau = 3$ 

ID	Strings
$s_1$	vankatesh
$s_2$	avataresha
$s_3$	kaushic chaduri
$S_4$	kaushik chakrab
$s_5$	kaushuk chadhui
$s_6$	caushik chakrabar

```
ED(s_1, s_2)=5 ED(s_1, s_3)=13 ED(s_1, s_4)=12 ED(s_1, s_5)=12 ED(s_1, s_6)=14 ED(s_2, s_3)=12 ED(s_2, s_4)=12 ED(s_2, s_5)=12 ED(s_2, s_6)=14 ED(s_3, s_4)=5 ED(s_3, s_5)=4 ED(s_3, s_6)=8 ED(s_4, s_5)=4 ED(s_4, s_6)=3 ED(s_5, s_6)=8
```

# **Data Cleaning & Integration**

#### **Relation with Duplicates**

ID	name	ZIP	Income
P1	Green	51519	30k
P2	Green	51518	32k
P3	Peter	30528	40k
P4	Peter	30528	40k
P5	Gree	51519	55k
P6	Chuck	51519	30k

### Challenges

 $O(n^2)$  pairs of strings. 1 million strings result in 1 trillion pairs!

O(|r||s|) time to calculate ED(r, s)

#### Filter-and-Refine Framework

- Basic idea
  - Filter a large number of dissimilar string pairs
  - Verify the remaining potentially similar pairs

- Good Filter Condition
  - Efficient to check
  - Effective for pruning dissimilar pairs

### Length Filter

• ED(r, s): The minimum number of edit operations (insertion/deletion/substitution) needed to transform r to s.

- Property:  $ED(r, s) \ge |r|-|s|$ 
  - it needs at least ||r|-|s|| deletions to just make r and s have the same length

#### **Applying the Length Filter**

- Give threshold  $\tau = 3$
- Pruning Condition:

$$\left| |s_i| - |s_j| \right| > 3$$

	\ /	
ID	Strings	Length
$s_1$	vankatesh	9
$s_2$	avataresha	10
$s_3$	kaushic chaduri	15
$s_4$	kaushik chakrab	15
$s_5$	kaushuk chadhui	15
$s_6$	caushik chakrabar	17

ED(
$$s_1$$
,  $s_2$ )=5 ED( $s_1$ ,  $s_3$ )=13 ED( $s_1$ ,  $s_4$ )=12 ED( $s_1$ ,  $s_5$ )=12 ED( $s_1$ ,  $s_6$ )=14 ED( $s_2$ ,  $s_3$ )=12 ED( $s_2$ ,  $s_4$ )=12 ED( $s_2$ ,  $s_5$ )=12 ED( $s_2$ ,  $s_5$ )=14 ED( $s_3$ ,  $s_4$ )=5 ED( $s_3$ ,  $s_5$ )=4 ED( $s_3$ ,  $s_6$ )=8 ED( $s_4$ ,  $s_5$ )=4 ED( $s_4$ ,  $s_6$ )=3 ED( $s_5$ ,  $s_6$ )=8

### **Partitioning Filter**

• Give threshold  $\tau = 1$ 



hi dose not appear in huston and needs at least 1 edit

### **Partitioning Filter**

• Give threshold  $\tau = 1$ 



minimum # of edit operations is 2. Prune!

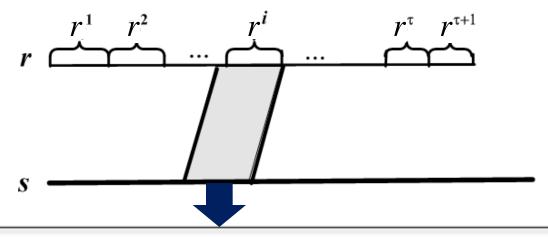
### **Partitioning Filter**

• Threshold au

split r to  $\tau + 1$  disjoint segments

• String r

• String s



Is there any substring of s matching a segment of r?



No

<*r*, *s*> are a candidate pair

<r, s> are dissimilar, prune

#### **How to Partition?**

• Give threshold  $\tau = 1$ 



Match



Candidate!

#### **Partition Scheme**

- Even Partition Scheme
  - Given  $\tau = 3$ , "avataresha"  $\rightarrow \{$  "av", "at", "are", "sha" $\}$

- Other Schemes
  - Select good partition strategies.
  - Adaptive partition scheme [Deng et al. 2012a].

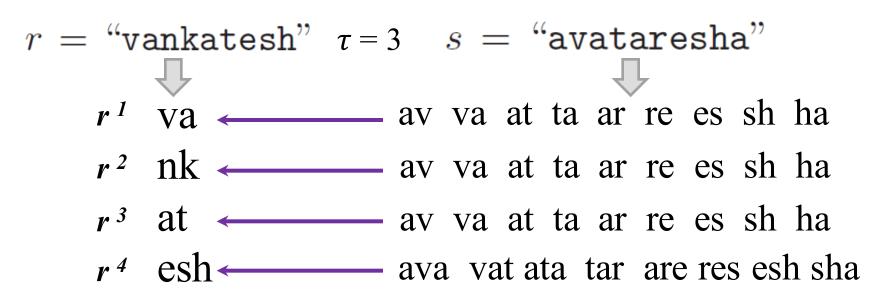
### Challenge

there are  $|s|^2$  substrings in s

how to reduce the number of substrings to compare with for each segment?

### **Length-based Method**

• For each segment, only compare to the substrings with the same length as the segment.



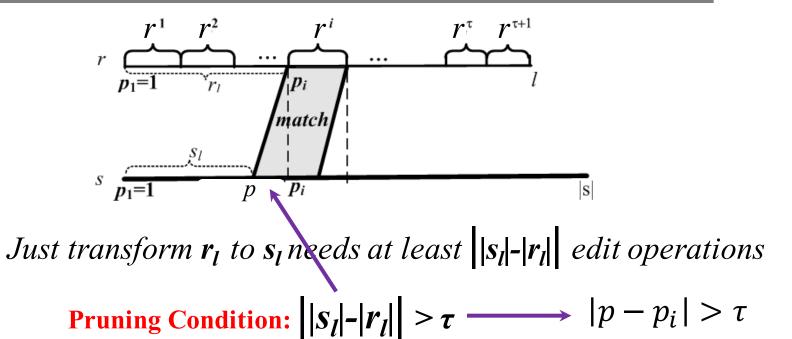
### **Length-based Method**

• For any strings r, s, and  $\tau$ , the number of comparisons:

$$(\tau+1)(|s|+1)-|r|$$

• For r = "vankatesh" and s = "avataresha", the number is 35

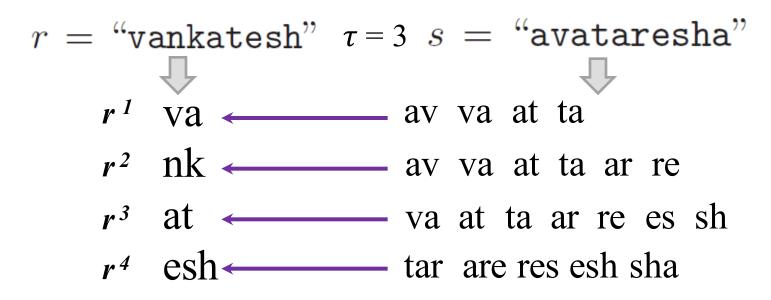
#### **Shift-based Method**



• For each segment  $r^i$  with the start position  $p_i$ , only compares to the substrings with start positions p in  $[p_i - \tau, p_i + \tau]$ 

#### **Shift-based Method**

• For each segment  $r^i$  with the start position  $p_i$ , only compares to the substrings with start positions p in  $[p_i - \tau, p_i + \tau]$ 



#### **Shift-based Method**

• For any strings r, s, and  $\tau$ , the number of comparisons:

$$(\tau + 1)(2\tau + 1)$$

• For r = "vankatesh" and s = "avataresha", the number is 22.

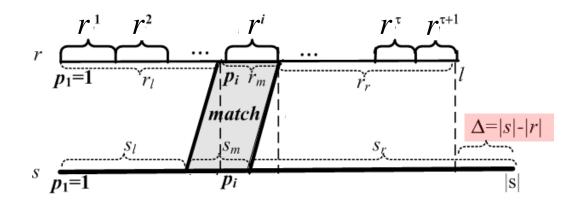
#### **Position-aware Method**

$$r= ext{"vankatesh"} \Longrightarrow \{ ext{va, nk, at esh}\}$$
 $s= ext{"avataresha"}$ 
 $s_l s_r$ 

Transform  $\mathbf{r_l}$  to  $\mathbf{s_l}$ , match "at", and transform  $\mathbf{r_r}$  to  $\mathbf{s_r}$ 

$$||s_l| - |r_l|| + ||s_r| - |r_r|| = 2 + 3 > \tau = 3$$

#### **Position-aware Method**



Transform  $\mathbf{r}_l$  to  $\mathbf{s}_l$ , match  $\mathbf{r}_m$  and  $\mathbf{s}_m$  then transform  $\mathbf{r}_r$  to  $\mathbf{s}_r$ 

**Pruning Condition:** 
$$||s_l|-|r_l||+||s_r|-|r_r||>\tau$$

• For each segment  $r^i$  with the start position  $p_i$ , only compare to the substrings with start position in  $[P_i - \left\lfloor \frac{\tau - \Delta}{2} \right\rfloor, P_i + \left\lfloor \frac{\tau + \Delta}{2} \right\rfloor]$  where  $\Delta = |s| - |r|$ 

### **Position-aware Method**

• For each segment  $r^i$  with the start position  $p_i$ , only compare to the substrings with start position in  $[P_i - \left\lfloor \frac{\tau - \Delta}{2} \right\rfloor, P_i + \left\lfloor \frac{\tau + \Delta}{2} \right\rfloor]$  where  $\Delta = |s| - |r|$ 

### **Position-aware Method**

• For any strings r, s, and  $\tau$ , the number of comparisons:

$$(\tau + 1)^2$$

• For r = "vankatesh" and s = "avataresha", the number is 14.

#### -- Left-side Perspective

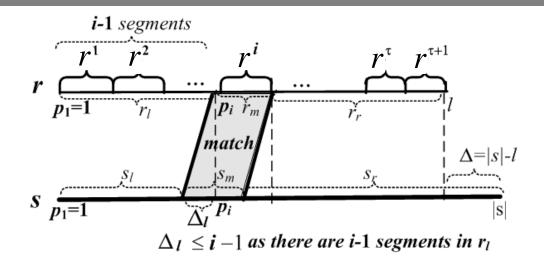
 $=4>\tau$ 

$$r=$$
 "vankatesh"  $\Rightarrow$  {va, nk, at, esh}  $r_l=$ "  $\tau=3$  3 unvisited segments  $s=$  "avataresha"  $s_l=$ "  $s_l=$ "  $s_l-|r_l|=1$ 

Thus we can safely skip the current matching segment and look for the next

matching substring in s, then r and s cannot be similar

#### -- Left-side Perspective



Pruning Condition:  $||s_l|-|r_l||+$  (# of unvisited segments) >  $\tau$ 

• For each segment  $r^i$  with the start position  $p_i$ , only compare to the substrings with start position in  $[P_i - (i-1), P_i + (i-1)]$ 

#### -- Left-side Perspective

• For each segment  $r^i$  with the start position  $p_i$ , only compare to the substrings with start position in  $[P_i - (i - 1), P_i + (i - 1)]$ 

$$\tau = 3$$
  $r =$  "vankatesh"  $s =$  "avataresha"

 $r^1$  Va  $\longleftarrow$  av

 $r^2$  nk  $\longleftarrow$  va at ta

 $r^3$  at  $\longleftarrow$  at ta ar re es

 $r^4$  esh  $\longleftarrow$  tar are res esh sha

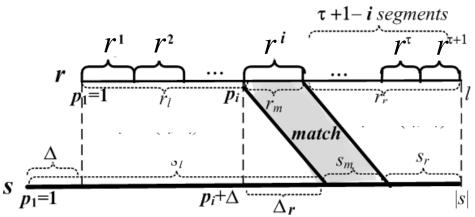
#### -- Left-side Perspective

• For any strings r, s, and  $\tau$ , the number of comparisons:

$$\tau^2 + 2\tau$$

• For r = "vankatesh" and s = "avataresha", the number is 14.

#### -- Right-side Perspective



 $\Delta_r \leq \tau + 1 - i$  as there are  $\tau + 1 - i$  segments in  $r_r$ 

## Pruning Condition: (# unvisited segments)+ $|s_r|-|r_r| > \tau$

• For each segment  $r^i$  with the start position  $p_i$ , only compare to the substrings with start position in  $[P_i + \Delta - (\tau + 1 - i), P_i + \Delta + (\tau + 1 - i)]$ 

• Interestingly, we can apply the multi-match-aware method from left- and right-side perspectives simultaneously.

• For each segment  $r^i$  with start position  $p_{i}$ , only compare to the substrings with start position in

$$[max(P_i - (i - 1), P_i + \Delta - (\tau + 1 - i)), \\ min(P_i + (i - 1), P_i + \Delta + (\tau + 1 - i))]$$

• For each segment  $r^i$  with start position  $p_i$ , only compare to the substrings with start position in  $[\max(P_i - (i-1), P_i + \Delta - (\tau + 1 - i)), \min(P_i + (i-1), P_i + \Delta + (\tau + 1 - i))]$ 

• For any strings r, s, and  $\tau$ , the number of comparisons:

$$\left[\frac{\tau^2 - \Delta^2}{2}\right] + \tau + 1$$

• For r = "vankatesh" and s = "avataresha", the number is 8.

### **Theoretical Results**

• The number of comparisons by the multi-match-aware method is **minimum** while guarantees completeness

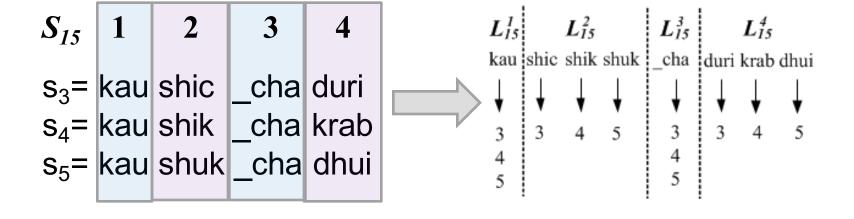
• For any s, r and  $\tau$ ,

$$W_{multi-match}(s, r, \tau) \subseteq W_{position}(s, r, \tau) \subseteq W_{shift}(s, r, \tau) \subseteq W_{length}(s, r, \tau)$$

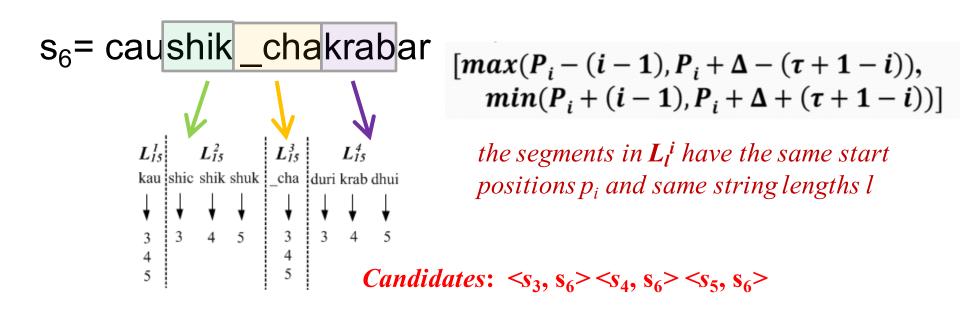
• 1. Group all the strings by length: S

ID	Strings	Length	
$s_1$	vankatesh	9	$S_9$
$s_2$	avataresha	10	$S_{10}$
$s_3$	kaushic chaduri	15	
$s_4$	kaushik chakrab	15	$S_{15}$
$s_5$	kaushuk chadhui	15	
$s_6$	caushik chakrabar	17	$S_{17}$

• 2. For each group  $S_{l}$ , partition its strings into  $\tau + 1$  segments and build  $\tau + 1$  inverted indexes  $L_{l}^{i}$ 



• 3. For each string s and index  $L_l^i$ , select substrings from s based on the partitioning filter to get candidates:



• 4. Verify the candidates

Candidates: 
$$< s_3, s_6 > < s_4, s_6 > < s_5, s_6 >$$

ED(
$$s_3, s_6$$
) > 3 X  
ED( $s_4, s_6$ ) = 3 X  
ED( $s_5, s_6$ ) > 3 X

# Improving Verification

# **Calculating Edit Distance**

	\$	b	r	i	t	n	e	y
\$	0	1	2	3	4	5	6	7
b	1	0	1	2	3	4	5	6
t	2	1	1	2	3	4	5	6
i	3	2	2	1	2	3	4	5
n	4	3	3	2	2	2	3	4
e	5	4	4	3	3	3	2	3
y	6	5	5	4	4	4	3	2

## **Verification**

	\$	0	1	2	3	4	5	6	7
	b	1	0	1	2	3	4	5	6
	t	2	1	1	2	3	4	5	6
ED(r[1 i] c[1 i])	i	3	2	2	1	2	3	4	5
$ED(r[1,i],s[1,j])$ $\geq  i-j $	n	4	3	3	2	2	2	3	4
— I* JI	e	5	4	4	3	3	3	2	3
	y	6	5	5	4	4	4	3	2

b

i

n

e

# **Verification**

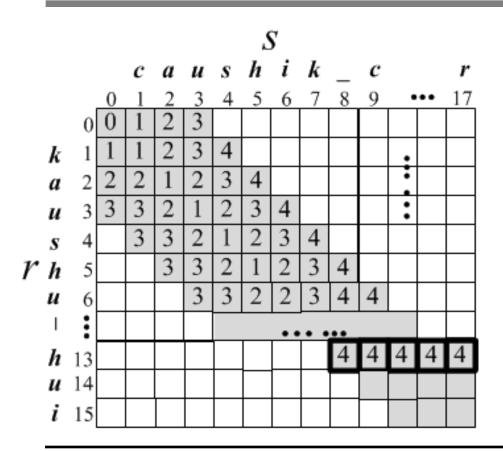
	\$	b	r	i	t	n	e	y
\$	0	1	2					
b	1	0	1	2				
t	2	1	1	2	3			
i		2	2	1	2	3		
n			3	2	2	2	3	
e				3	3	3	2	3
y					4	4	3	2

## Verification

only need to calculate a band of width  $2\tau+1$ 

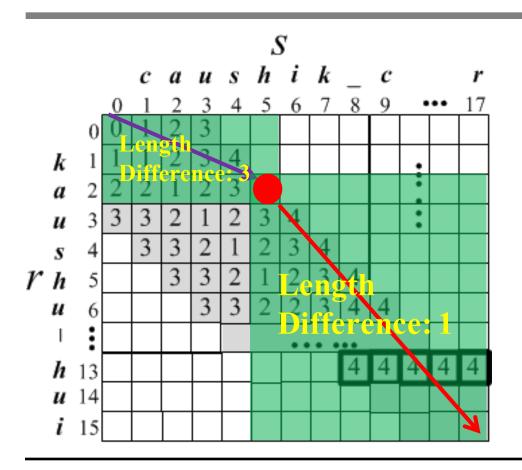
		\$	b	r	i	t	n	e	y
	\$	0	1	2					
	b	1	0	1	2				
$\ell$	t	2	1	1	2	3			
	i		2	2	1	2	3		
	n			3	2	2	2	3	
	e				3	3	3	2	3
	y					4	4	3	2

## **Early Termination**



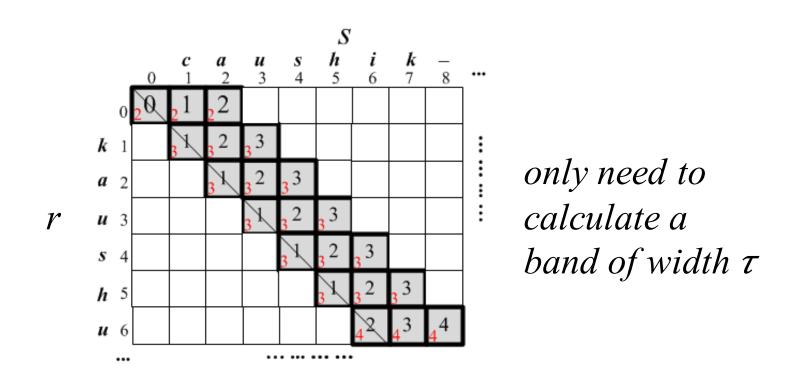
all derived cells must have values larger than  $\tau$ .

## Length-aware Verification

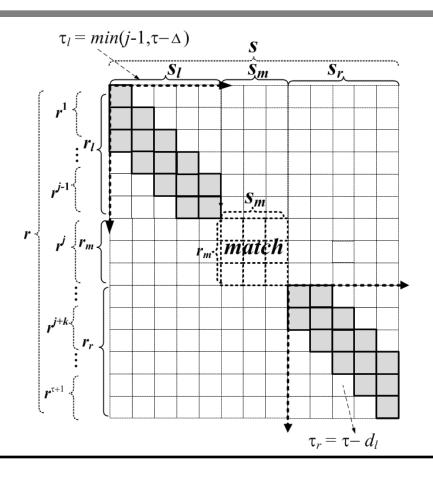


total length difference is  $4 > \tau$ . Thus no need to calculate M[2][5].

## Length-aware Verification



## **Extension-base Verification**



### **Extension-base Verification**

- We can verify a candidate pair using tighter thresholds:
  - For the left parts we can set  $\tau_l = i 1$ .
  - For the right parts we can set  $\tau_r = \tau + 1 i$ .

band widths are  $\tau_l < \tau$  and  $\tau_r < \tau$ 

## **Takeaways**

- (1) The partitioning filter for Edit Distance
- (2) The multi-match-aware method
- (3) The extension-based verification

### References

- Pass-Join: A Partition based Method for Similarity Joins. G. Li, D. Deng, J. Wang, J. Feng. VLDB 2012.
- A Pivotal Prefix Based Filtering Algorithm for String Similarity Search. D. Deng, G. Li, J. Feng. SIGMOD 2014.
- Ed-Join: An Efficient Algorithm for Similarity Joins with Edit Distance Constraints. C. Xiao, W. Wang, X. Lin. VLDB 2008