

## Introduction

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# Introduction

# COMP90049 **Knowledge Technologies**

Jeremy Nicholson and Justin Zobel, Karin Verspoor and Rao Kotagiri

Semester 1





# Summary

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#### Neighbo

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## Week 2:

- Approximate String Search and Matching
- Common Applications
- Methods:
  - Neighbourhood Search
  - Edit Distance
  - N-Gram Distance
  - Phonetic methods
- Evaluation



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## Consider:

- Given a string, is some substring contained within it?
- Given a string (document), find all occurrences of some substring



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## For example, find Exxon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.



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## For example, find Exxon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo  $\bf Exxon$  max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.



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## Consider:

- Given a string, is some substring contained within it?
- Given a string (document), find all occurrences of some substring

Not (really) a Knowledge Technology!



# Approximate String Search

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# String Sean Exact

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### Find exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.



# Approximate String Search

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## Find exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.

# Not present!

...But what is the "closest" or "best" match?



# Approximate String Search

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# Approximate

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## Find exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.

## Not present!

...But what is the "closest" or "best" match?

This is a Knowledge Technology!



# Important problems

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# Two main applications for Approximate String Search:

- Spelling correction
- Computational Genomics



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# Need the notion of a **dictionary**:

Here, a list of words



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# Need the notion of a dictionary:

 Here, a list of words entries that are "correct" with respect to our (expectations of our) language



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- Here, a list of words entries that are "correct"
- We can break our input into words substrings that we wish to match, and compare each of them against the entries in the dictionary



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- Here, a list of words entries that are "correct"
- We can break our input into words substrings that we wish to match, and compare each of them against the entries in the dictionary
- A word item in the input which doesn't appear in the dictionary is misspelled



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- Here, a list of words entries that are "correct"
- We can break our input into words substrings that we wish to match, and compare each of them against the entries in the dictionary
- A word item in the input which doesn't appear in the dictionary is misspelled
- A werd item in the input which does appear in the dictionary might be correctly spelled or misspelled



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- Here, a list of words entries that are "correct"
- We can break our input into words substrings that we wish to match, and compare each of them against the entries in the dictionary
- A word item in the input which doesn't appear in the dictionary is misspelled
- A word item in the input which does appear in the dictionary might be correctly spelled or misspelled (probably slightly beyond the scope of this subject)



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Therefore, the problem here:

Given some item of interest — which does not appear in our dictionary — which entry from the dictionary was truly intended?



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Therefore, the problem here:

Given some item of interest — which does not appear in our dictionary — which entry from the dictionary was truly intended?

Depends on the person who wrote the original string!



# Computational Genomics

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## Typical Genomics problem:

- Given a nucleotide/amino acid sequence (substring)
- Find whether the sequence occurs within a larger sequence (string)
- Possibly with "errors" (nucleotide/amino acid changes)



# Computational Genomics

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## Typical Genomics problem:

- Given a substring, find whether the sequence occurs within a larger string, possibly with "errors"
- Almost the same problem, flipped around



# **Computational Genomics**

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# Typical Genomics problem:

- Given a substring, find whether the sequence occurs within a larger string, possibly with "errors"
- Almost the same problem
- But much larger strings: a small genomics problem might involve comparing perhaps 1K character sequence against several 100K character sequences; alphabet is smaller



# Other Problems of Interest

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■ Name matching, for example:

The name *Gorbachev* is spelled (at least) 20 different ways in a corpus of newswire text!

Gorbachev, Gorbachev, Gorbahev, Gorbatchev, Gorbechev, Gorbachov, Gorachev, Gorbacheva, Gorbachev, Gorbachev, Gorbachev, Gorbachev, ...



# Other Problems of Interest

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Name matching

- Query repair
- Phonetic matching
- Data cleaning
- ...



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Find approximate match(es) for exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.



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Find approximate match(es) for exon in:

In exes for foxes rex dux mixes a pox of waxed luxes.

An axe, and an axon, to exo **Exxon** max oxen.

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Insert x (and fold case)



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Find approximate match(es) for exon in:

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Delete n



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Find approximate match(es) for exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an **axon**, to exo Exxon max oxen. Grexit or Brexit as quixotic haxxers with buxom rex taxation.

Replace e with a (Sometimes Substitute)



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Find approximate match(es) for exon in:

In exes for foxes rex dux mixes a pox of waxed luxes. An axe, and an axon, to exo Exxon max **oxen**. Grexit or Brexit as quixotic haxxers with buxom rex taxation.

**Transpose** e and o (Beyond the scope of this subject.)



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For a given string w of interest:



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# For a given string *w* of interest:

 Generate all variants of w that utilise at most k changes (Insertions/Deletions/Replacements) — neighbours



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# For a given string w of interest:

- Generate all variants of w that utilise at most k changes (Insertions/Deletions/Replacements) — neighbours
- Check whether generated variants exist in dictionary



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## For a given string *w* of interest:

- Generate all variants of w that utilise at most k changes (Insertions/Deletions/Replacements) — neighbours
- Check whether generated variants exist in dictionary
- All results found in dictionary are returned

Unix command-line utility agrep is an efficient mechanism for finding these.



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## For example:

... proceed if you can see no ther option ...



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## For example:

... proceed if you can see no **ther** option ...

Intended word: other

Requires 1 insertion (o) so intended word will be found using neighbourhood search (and some unintended words...)



# Neighbourhood Search Efficiency

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With a careful implementation, Neighbourhood search is suprisingly fast!



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Neighbourhood search is suprisingly fast!

Consider: alphabet size is  $\Sigma$ , length of string is |w|:

For 1 edit, roughly  $\mathcal{O}(\Sigma \cdot |w|)$  neighbours



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Neighbourhood search is suprisingly fast!

Consider: alphabet size is  $\Sigma$ , length of string is |w|:

For 2 edits, roughly  $\mathcal{O}(\Sigma^2 \cdot |w|^2)$  neighbours



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Neighbourhood search is suprisingly fast!

Consider: alphabet size is  $\Sigma$ , length of string is |w|:

For k edits, roughly  $\mathcal{O}(\Sigma^k \cdot |w|^k)$  neighbours



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Neighbourhood search is suprisingly fast!

Consider: alphabet size is  $\Sigma$ , length of string is |w|:

...But  $\Sigma$  is a small constant, string of interest is usually short, and k is usually small

For k edits, roughly  $\mathcal{O}(\Sigma^k \cdot |w|^k)$  neighbours



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Neighbourhood search is suprisingly fast!

Consider: alphabet size is  $\Sigma$ , length of string is |w|:

For k edits, roughly  $\mathcal{O}(\Sigma^k \cdot |w|^k)$  neighbours

For each neighbour, need a dictionary read (dict has D entries): In total,  $\mathcal{O}(\Sigma^k \cdot |w|^k \log D)$  string comparisons

...But  $\Sigma$  is a small constant, string of interest is usually short, and k is usually small

For each neighbour, need a dictionary read (dict has D entries): In total,  $\mathcal{O}(|w|^k \log D)$  string comparisons



## Neighbourhood Search Effectiveness

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So, efficiency isn't our problem.

 $({\tt agrep}\ example)$ 



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Alternative method:

Scan through each dictionary entry looking for the "best" match



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Evaluation References Global Edit Distance:

Transform the string of interest into each dictionary entry, using the operations Insert, Delete, Replace, and Match (character)



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### Global Edit Distance:

Transform the string of interest into each dictionary entry, using the operations Insert, Delete, Replace, and Match (character)

Each operation is associated with a score; Best match is the dictionary entry with best aggregate score



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## Edit Distance

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For example:

Item of interest: crat

Dictionary: cart, arts



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For example:

Item of interest: crat

Dictionary: cart, arts

 $\mathtt{crat} \to \mathtt{cart}$ :

Match c, Delete r, Match a, Insert r, Match t



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**Edit Distance** 

For example:

Item of interest: crat

Dictionary: cart, arts

 $crat \rightarrow cart$ :

Match c, Delete r, Match a, Insert r, Match t

 $crat \rightarrow arts$ :

Replace c with a, Match r, Delete a, Match t, Insert s



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**Edit Distance** 

References

For example:

Item of interest: crat

Dictionary: cart, arts

Score: Match +1, Insert -1, Delete -1, Replace -1

 $\mathtt{crat} \to \mathtt{cart}$ :

Match c, Delete r, Match a, Insert r, Match t

 $\mathtt{crat} \to \mathtt{arts}$ :

Replace c with a, Match r, Delete a, Match t, Insert s



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**Edit Distance** 

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For example:

Item of interest: crat

Dictionary: cart, arts

Score: Match +1, Insert -1, Delete -1, Replace -1

 $crat \rightarrow cart$ :

Match c (+1), Delete r (-1), Match a (+1), Insert r (-1), Match t (+1) = +1

 $crat \rightarrow arts$ :

Replace c with a (-1), Match r (+1), Delete a (-1), Match t (+1), Insert s (-1) = -1

cart, is the better match



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Confusingly, Global Edit Distance isn't a "distance"



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Confusingly, Global Edit Distance isn't a "distance"

...But depends on parameter



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Match (0), Insert (+1), Delete (+1), Replace (+1)

This is the Levenshtein Distance (true distance): number of edits required to transform one string into the other (symmetric)



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Reference:

Hypothetically, any parameter is possible!



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Hypothetically, any parameter is possible!

But some choices make no sense, e.g.:

Match (+4), Insert (-2), Delete (+8), Replace (0)



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Hypothetically, any parameter is possible!

But some choices make no sense, e.g.:

Match (+4), Insert (-2), Delete (+8), Replace (0)

Which corresponds to best match?

- Insert, Delete, Insert, Delete, Insert, Delete
- Match, Match, Match
- Replace, Match, Replace



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Hypothetically, any parameter is possible!

But some choices make no sense, e.g.:

Match (+4), Insert (-2), Delete (+8), Replace (0)

Which corresponds to best match?

- Insert, Delete, Insert, Delete = +18
- Match, Match, Match = +12
- Replace, Match, Replace = +4



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Often, "direction" doesn't matter: Insert = Delete ("Indel")



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Sometimes, score of Replace depends on which character is being replaced:

Consider:

Is faxing more likely to be facing or faking?



## Global Edit Distance Algorithm

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Computer can't find best sequence of operations by inspection



## Global Edit Distance Algorithm

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From string f to string t, given array of size |f|+1 by |t|+1, we can solve using the Needleman–Wunsch algorithm:



# Global Edit Distance Algorithm

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From string t, given array A of size |t|+1 by |t|+1, we can solve using the Needleman–Wunsch algorithm:

```
i = Insertion cost
d= Deletion cost
equal() returns m if characters match, r otherwise
lf = strlen(f); lt = strlen(t);
A[0][0]=0:
for (j=1; j<=1t; j++) A[j][0] = j * i;
for (k=1; k<=1f; k++) A[0][k] = k * d;
for (j=1; j<=lt; j++)
   for (k=1; k<=lf; k++)
      A[j][k] = max3( //Or min3 if m<i,d,r
         A[i][k-1] + d, //Deletion
         A[j-1][k] + i, //Insertion
         A[j-1][k-1] + equal(f[k-1],t[j-1])); %Replace or match
```

Final score is at A[lt][lf]



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	ε	С	r	a	t
ε					
a					
r					
t					
s					



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

Initialise table:

	ε	С	r	a	t
$\overline{\varepsilon}$	0	-1	-2	-3	-4
a	-1				
r	-2				
t	-3		-2		
s	-4				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, consider three neighbours:

	ε	С	r	a	t
ε	0	-1	-2	-3	-4
a	-1	?	-2		
r	-2				
t	-3				
s	-4				



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For c-a correspondence, Delete c:

	ε		r		t
$\varepsilon$	0 -1 -2 -3	-1	-2	-3	-4
a	-1	-2			
r	-2				
t	-3				
s	-4				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, Insert a:

	ε	С	r	a	t
$\varepsilon$	0	-1	-2	-3	-4
	-1	-2	-2		
r t	-1 -2 -3				
t	-3				
s	-4				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, Replace c with a:

		С		a	t
ε	0	-1	-2	-3	-4
a	-1	-1	-2		
r	-2				
t	-3				
s	-4				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

			r		t
$\varepsilon$	0	-1	-2	-3	-4
a	-1	-1	-2		
r	-2				
t	-3		-2 -2		
s	-4				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

	ε	С	r	a	t
$\varepsilon$	0	-1	-2	-3	-4
a	-1	-1	-2	-1	
r	-2		- <mark>2</mark> -2		
t	-3				
s	-4				



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Evaluatio

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

	$\varepsilon$	С	r	a	t
$\overline{\varepsilon}$	0	-1	-2	-3	-4
a	-1	-1	-2	-1	-2
r	-2	-2	0	-3 -1 -1 -1 -2	-2
t	-3	-3	-1	-1	0
s	-4	-4	-2	-2	-1



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References

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

	$\varepsilon$	С	r	a	t
ε	0	-1	-2	-3	-4
a	-1	-1 -1 -2 -3 -4	-2	-1	-2
r	-2	-2	0	-1	-2
t	-3	-3	-1	-1	0
s	-4	-4	-2	-2	-1

Global Edit Distance: -1 (Replace, Match, Delete, Match, Insert)



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#### String Seam

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Dhonotio

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Algorithm actually depends on parameter!



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```
A[j][k] = max3(
   A[j][k-1] + d, //Deletion
   A[j-1][k] + i, //Insertion
   A[j-1][k-1] + equal(f[k-1],t[j-1])); //Replace or match
```



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```
A[j][k] = max3(
    A[j][k-1] + d, //Deletion
    A[j-1][k] + i, //Insertion
    A[j-1][k-1] + equal(f[k-1],t[j-1])); //Replace or match
```

→ Match score greater than Insert/Delete/Replace

```
e.g. Match (+1), Insert/Delete/Replace (-1)
```



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A[j][k] = min3(
 A[j][k-1] + d, //Deletion
 A[j-1][k] + i, //Insertion
 A[j-1][k-1] + equal(f[k-1],t[j-1])); //Replace or match



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```
A[j][k] = min3(
    A[j][k-1] + d, //Deletion
    A[j-1][k] + i, //Insertion
    A[j-1][k-1] + equal(f[k-1],t[j-1])); //Replace or match
```

→ Match score less than Insert/Delete/Replace

```
e.g. Match (0), Insert/Delete/Replace (+1)
```

(Levenshtein Distance)



# Local Edit Distance

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Local Edit Distance is like Global Edit Distance, but we are searching for the best <u>substring</u> match



# **Local Edit Distance**

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Local Edit Distance is like Global Edit Distance, but we are searching for the best substring match

Particularly suitable when comparing two strings of very different lengths, e.g. a word and a sentence



# Local Edit Distance Algorithm

lf = strlen(f); lt = strlen(t);

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. ......

References

From string f to string t, given array A of size |f|+1 by |t|+1, we can solve using the Smith–Waterman algorithm:

equal() returns m if characters match, r otherwise

Final score is greatest value in the entire table (or least value, if m < i, d, r)



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N-Grani Disi

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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

(For Local Edit Distance, Match  $\underline{\text{must}}$  have different +/- sign to Insert/Delete/Replace)



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DI. . . . . . . . . .

\_ . . .

References

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)  $\,$ 

	ε	С	r	a	t
$\varepsilon$					
a					
r					
t					
s					



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

Initialise table:

	$\varepsilon$	С	r	a	t
$\overline{\varepsilon}$	0 0 0 0	0	0	0	0
a	0				
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, consider three neighbours:

	$\varepsilon$		r	a	t
$\varepsilon$	0	0 ?	0	0	0
arepsilon arts	00000	?			
r	0				
t	0				
s	0				



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References

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, Delete c:

	$\varepsilon$	С	r	a	t
ε	0	0 -1	0	0	0
arepsilon arts	0 0 0 0	-1			
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, Insert a:

	$\varepsilon$	С	r	a	t
ε	0	0	0	0	0
$\varepsilon$ a r t	0 0 0 0	-1			
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, Replace c with a:

	$\varepsilon$	С	r	a	t
ε	0	0 -1	0	0	0
$\varepsilon$ a r t	0 0 0	-1			
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For c-a correspondence, 0 is better:

	ε	С	r	a	t
ε	0	0	0	0	0
a	0	0		0	
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For r-a correspondence, 0 is better:

	$\varepsilon$	С	r	a	t
$\varepsilon$	0	0	0	0	0
a	0	0 0	0		
r	0				
t	0				
s	0				



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In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

For a-a correspondence, Match:

	ε		r		t
$\varepsilon$	0	0	0	0	0
arepsilon a	0	0	0	1	
r	0				
t	0				
s	0				



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Evaluation

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

	ε	С	r	a	t
$\overline{\varepsilon}$	0	0	0	0	0
a	0	0	0	1	0
r	0	0	1	0	0
t	0	0	0	0	1
s	0	0	0	0 1 0 0	0



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**Edit Distance** 

References

In action: from crat to arts, Match (+1), Insert/Delete/Replace (-1)

And so on:

7110 30 011.

	ε	С	r	a	t
$\varepsilon$	0	0	0	0	0
a	0	0	0	1	0
r	0	0	1	0	0
t	0	0	0	0	1
s	0 0 0 0	0	0	0	0

Three (equivalent) subsequences tied for best match (+1)



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For strings f and t, Both algorithms above are  $\mathcal{O}(|f||t|)$  in both space and time.



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Evaluation

When approximate matching, we have a constant string f which we want to compare to <u>each</u> string in the dictionary:



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N-Gram Dis Conomico

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When approximate matching, we have a constant string f which we want to compare to each string t in the dictionary D:

$$\mathcal{O}(\sum_{t \in D} |f||t|)$$



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#### Genomics

\_ . . .

Evaluation

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When approximate matching, we have a constant string f which we want to compare to each string t in the dictionary D:

$$\mathcal{O}(|f|\sum_{t\in D}|t|)$$



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When approximate matching, we have a constant string f which we want to compare to each string t in the dictionary D:

Hence, integer comparisons are roughly the number of characters in the dictionary. Whether this is feasible depends on the size of the dictionary.



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N-Gram Distance has same goal as Edit Distance: compare two strings to determine "best" match

A true "distance"



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N-Gram Distance has same goal as Global Edit Distance, but much simpler



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(character) n-gram: substring of length n



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n-gram: substring of length n

2-grams of crat: cr, ra, at



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n-gram: substring of length n

2-grams of crat: #c, cr, ra, at, t# (sometimes)



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Frankrickie

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n-gram: substring of length n

3-grams of crat: #cr, cra, rat, at#



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#### Diameter.

Contrasta.

References

n-gram: substring of length n

2-grams of crat: #c, cr, ra, at, t#

2-grams of cart: #c, ca, ar, rt, t#

2-grams of arts: #a, ar, rt, ts, s#



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*n*-gram: substring of length *n* 

2-grams of crat: #c, cr, ra, at, t#

2-grams of cart: #c, ca, ar, rt, t#

2-grams of arts: #a, ar, rt, ts, s#

N-Gram Distance between *n*-grams of string  $s(G_n(s))$  and  $t(G_n(t))$ :

$$|G_n(s)|+|G_n(t)|-2\times |G_n(s)\cap G_n(t)|$$

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N<sub>e</sub>Gram Distance

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*n*-gram: substring of length *n* 

2-grams of crat: #c, cr, ra, at, t#

2-grams of cart: #c, ca, ar, rt, t#

2-grams of arts: #a, ar, rt, ts, s#

2-Gram Distance between crat and cart:

 $|G_2(\mathtt{crat})| + |G_2(\mathtt{cart})| - 2 \times |G_2(\mathtt{crat}) \cap G_2(\mathtt{cart})|$ 

 $= 5 + 5 - 2 \times 2 = 6$ 

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N<sub>e</sub>Gram Distance

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*n*-gram: substring of length *n* 

2-grams of crat: #c, cr, ra, at, t#

2-grams of cart: #c, ca, ar, rt, t#

2-grams of arts: #a, ar, rt, ts, s#

2-Gram Distance between crat and cart:

$$|\textit{G}_{2}(\texttt{crat})| + |\textit{G}_{2}(\texttt{cart})| - 2 \times |\textit{G}_{2}(\texttt{crat}) \cap \textit{G}_{2}(\texttt{cart})|$$

$$=5+5-2\times 2=6$$

2-Gram Distance between crat and arts:

$$|G_2(\text{crat})| + |G_2(\text{arts})| - 2 \times |G_2(\text{crat}) \cap G_2(\text{arts})|$$

$$= 5 + 5 - 2 \times 0 = 10$$

# N-Gram Distance

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*n*-gram: substring of length *n* 

2-grams of crat: #c, cr, ra, at, t#

2-grams of cart: #c, ca, ar, rt, t#

2-grams of arts: #a, ar, rt, ts, s#

2-Gram Distance between crat and cart:

$$|G_2(\operatorname{crat})| + |G_2(\operatorname{cart})| - 2 \times |G_2(\operatorname{crat}) \cap G_2(\operatorname{cart})|$$

$$= 5 + 5 - 2 \times 2 = 6$$
 (better)

2-Gram Distance between crat and arts:

$$|G_2(\text{crat})| + |G_2(\text{arts})| - 2 \times |G_2(\text{crat}) \cap G_2(\text{arts})|$$
  
= 5 + 5 - 2 × 0 = 10



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Doforonco

Occasionally useful as a simpler variant of (Global) Edit Distance



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Evaluation References Occasionally useful as a simpler variant of Edit Distance

More sensitive to long substring matches, less sensitive to relative ordering of strings (matches can be anywhere!)



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References

Occasionally useful as a simpler variant of Edit Distance

More sensitive to long substring matches, less sensitive to relative ordering of strings (matches can be anywhere!)

Despite its simplicity, takes roughly the same time to compare entire dictionary



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Occasionally useful as a simpler variant of Edit Distance

More sensitive to long substring matches, less sensitive to relative ordering of strings (matches can be anywhere!)

Despite its simplicity, takes roughly the same time to compare entire dictionary

Quite useless for very long strings and/or very small alphabets (Why?)



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## Genomics

\_ . . .

References

Recall: we have a "short" ( $\sim$ 1K character) nucleotide/amino acid sequence to compare against many long ( $\sim$ 100K character) chromosomes/genes/proteins/etc.



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# Neighbourl

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#### Genomics

Reference

Recall: we have a "short" ( $\sim$ 1K character) string to compare against many long ( $\sim$ 100K character) strings



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Recall: we have a "short" ( $\sim$ 1K character) string to compare against many long ( $\sim$ 100K character) strings

For example, if some member of the population has 99% of the sequence of interest, they might be susceptible to some medical condition



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## Edit Distan

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References

Recall: we have a "short" ( $\sim$ 1K character) string to compare against many long ( $\sim$ 100K character) strings

We're allowed  $\sim$ 10 errors; alphabet is  $\sim$ 4 or  $\sim$ 20 characters



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Neighbourhood search:

Roughly  $4^{10}\times 1000^{10}$  possible neighbours.



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#### Mojekh

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Neighbourhood search:

Roughly  $4^{10} \times 1000^{10}$  possible neighbours.

... Forget it.



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#### Najahhau

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## Global Edit Distance:

One string is  ${\sim}1K$  characters, other is  ${\sim}100K$  characters.



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Global Edit Distance:

One string is  $\sim 1 \text{K}$  characters, other is  $\sim 100 \text{K}$  characters.

Complexity  $\sim 1K * 100K = 10^3 * 10^5 = 10^8$ 

→ Prefers shorter chromosomes (not intended behaviour)



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## Local Edit Distance:

One string is  $\sim$ 1K characters, other is  $\sim$ 100K characters.

... Seems like the right idea.



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Local Edit Distance:

One string is  $\sim 10 K$  characters, other is  $\sim 1 G$  characters.

... Can't fit table into memory.



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Local Edit Distance:

One string is  $\sim$ 10K characters, other is  $\sim$ 1G characters.

... Requires approximate solutions with heuristics, e.g. BLAST, FASTA



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## N-Gram Distance:

With huge *n* (e.g. 80% of length of shorter string) can (almost) work!



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N-Gram Distance:

Surprisingly, can (almost) work!

Tends to prefer shorter chromosomes like Global Edit Distance



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## N-Gram Distance:

But better methods for using *n*-gram information, e.g. de Bruijn graphs



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# Genomics

## **Phonetics**

Poforonco

In English (and some other languages), **orthography** (spelling) isn't a good predictor of **phonetics** (sounds)



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In English (and some other languages), **orthography** (spelling) isn't a good predictor of **phonetics** (sounds)

Salient concern in speech—to—text systems, e.g.: Georgia Conal



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In English (and some other languages), **orthography** (spelling) isn't a good predictor of **phonetics** (sounds)

Salient concern in speech—to—text systems, e.g.: Georgia Conal George O'Connell



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In English (and some other languages), **orthography** (spelling) isn't a good predictor of **phonetics** (sounds)

Salient concern in speech-to-text systems, e.g.: Lho, Lo, Loan, Loe, Loew, Lough, Low, Lowe, ...



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N-Gram Distance

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In English (and some other languages), **orthography** (spelling) isn't a good predictor of **phonetics** (sounds)

Also relevant in spelling correction (English can be very difficult to spell correctly!)



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#### **Phonetics**

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One (ineffectual) mechanism: Soundex



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One mechanism: Soundex

```
{\tt aehiouwy} \quad \rightarrow \quad 0 \; (vowels)
```

 ${\tt bpfv} \quad \to \quad {\tt 1} \; ({\tt labials})$ 

 $\texttt{cgjkqsxz} \quad \rightarrow \quad \textbf{2 (misc: fricatives, velars, etc.)}$ 

Translation table: dt  $\rightarrow$  3 (dentals)

 $1 \rightarrow 4$  (lateral)

mn  $\rightarrow$  5 (nasals)

 $r \rightarrow 6$  (rhotic)



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One mechanism: Soundex

aehiouwy  $\rightarrow$  0 (vowels) bpfv  $\rightarrow$  1 (labials)

 $\texttt{cgjkqsxz} \quad \rightarrow \quad \textbf{2 (misc: fricatives, velars, etc.)}$ 

Translation table:  $\hspace{1.5cm} \mathtt{dt} \hspace{0.5cm} \rightarrow \hspace{0.5cm} 3 \hspace{0.1cm} (\text{dentals})$ 

 $1 \rightarrow 4 \text{ (lateral)}$ 

mn  $\rightarrow$  5 (nasals)

 $r \rightarrow 6$  (rhotic)

#### **Phonetics**

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Four step process:

- Except for initial character, translate string characters according to table
- **2** Remove duplicates (e.g.  $4444 \rightarrow 4$ )
- Remove 0s
- Truncate to four symbols



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One mechanism: Soundex

 $\begin{array}{ccc} \texttt{aehiouwy} & \rightarrow & \texttt{0 (vowels)} \\ \texttt{bpfv} & \rightarrow & \texttt{1 (labials)} \end{array}$ 

 $\texttt{cgjkqsxz} \quad \rightarrow \quad \textbf{2 (misc: fricatives, velars, etc.)}$ 

Translation table: dt  $\rightarrow$  3 (dentals)

 $1 \rightarrow 4 \text{ (lateral)}$ mn  $\rightarrow 5 \text{ (nasals)}$ 

 $\texttt{r} \quad \rightarrow \quad \texttt{6 (rhotic)}$ 

N-Gram Distance

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Four step process:

king kyngge k052 k05220 k052 k0520 k52 k52



### Introduction

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One mechanism: Soundex

 $\begin{array}{ccc} \texttt{aehiouwy} & \rightarrow & \texttt{0 (vowels)} \\ \texttt{bpfv} & \rightarrow & \texttt{1 (labials)} \end{array}$ 

 $\texttt{cgjkqsxz} \quad \rightarrow \quad \textbf{2 (misc: fricatives, velars, etc.)}$ 

Translation table:  $dt \rightarrow 3 \text{ (dentals)}$ 

 $1 \rightarrow 4 \text{ (lateral)}$ mn  $\rightarrow 5 \text{ (nasals)}$ 

 $\texttt{r} \quad \rightarrow \quad \texttt{6 (rhotic)}$ 

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Four step process:

knight night k50203 n0203 k50203 n0203 k523 n23



### Introduction

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One mechanism: Soundex

 $\begin{array}{ccc} \texttt{aehiouwy} & \rightarrow & \texttt{0 (vowels)} \\ \texttt{bpfv} & \rightarrow & \texttt{1 (labials)} \end{array}$ 

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Translation table: dt  $\rightarrow$  3 (dentals)

 $\begin{array}{ccc} \textbf{1} & \rightarrow & \textbf{4 (lateral)} \\ \textbf{mn} & \rightarrow & \textbf{5 (nasals)} \end{array}$ 

 $\texttt{r} \quad \rightarrow \quad \textbf{6 (rhotic)}$ 

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# Four step process:

loan	loew	lough	lewicks
1005	1000	10020	1000222
105	10	1020	102
15	1	12	12



# Other Phonetic Methods

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Better phonetic methods make use of the fact that some letters sounds alike in certain contexts, and different in other contexts

**Editex** uses the Edit Distance to compare strings based on a similar translation table to Soundex

**Ipadist** uses a text–to–sound algorithm to represent tokens according to the International Phonetic Alphabet (but context matters a lot)

There are also worse variants, like Phonix.



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### **Evaluation**

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Evaluation: consider whether the system is effective at solving the user's problem



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Evaluation: consider whether the system is effective at solving the user's problem

In this case: for a misspelled word, does the system identify the correct word?



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To evaluate, we need:

A number of cases of misspelled words



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# To evaluate, we need:

- A number of cases of misspelled words
- The intended (correct) word for each case



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# To evaluate, we need:

- A number of cases of misspelled words
- The <u>intended</u> (correct) word for each case
- An evaluation metric



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### **Evaluation**

We have some cases:



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#### **Evaluation**

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Misspelled Word	Correct Word
ther	other
corridr	corridor
cracheyt	crotchety



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Misspelled Word	Correct Word	Predicted Word
ther	other	there
corridr	corridor	corridor
cracheyt	crotchety	cachet



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#### **Evaluation**

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Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
ther	other	there	×
corridr	corridor	corridor	✓
cracheyt	crotchety	cachet	×



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Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
ther	other	there	×
corridr	corridor	corridor	✓
cracheyt	crotchety	cachet	×

**Accuracy**: fraction of correct responses



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Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
ther	other	there	×
corridr	corridor	corridor	✓
cracheyt	crotchety	cachet	×

Accuracy: Number of correct predictions
Total number of words



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#### **Evaluation**

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## More realistic situation:

Misspelled Word	Correct Word	Predicted Word
		there
ther	other	ether
		their
corridr	corridor	corridor
Corridr	Corridor	carrier
cracheyt	crotchety	???
•••		



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### More realistic situation:

Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
		there	×
ther	other	ether	×
		their	×
corridr	corridor	corridor	✓
COLLIGI	COITIGOI	carrier	×
cracheyt	crotchety	???	???
•••			



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Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
		there	×
ther	other	ether	×
		their	×
corridr	corridor	corridor	✓
Collidi	Corridor	carrier	×
cracheyt	crotchety	???	_

Precision: fraction of correct responses among attempted responses



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Misspelled Word	Correct Word	Predicted Word	Right/Wrong?
		there	×
ther	other	ether	×
		their	×
corridr	corridor	corridor	✓
COLLIGI	COLLIGOL	carrier	×
cracheyt	crotchety	???	_

Recall: proportion of words with a correct response (somewhere)



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#### **Evaluation**

Typically, the value of the evaluation metric has little intrinsic meaning



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#### **Evaluation**

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Typically, the value of the evaluation metric has little intrinsic meaning

"This system gets 81% accuracy" — useful for users, or not?



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The evaluation metric allows us to <u>compare</u> systems:

"The system based on the Global Edit Distance gets 81% accuracy, whereas the system based on the N-Gram Distance gets 84% accuracy"



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The evaluation metric allows us to <u>compare</u> systems:

"The system based on the Global Edit Distance gets 81% accuracy, whereas the system based on the N-Gram Distance gets 84% accuracy" — Why?



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The evaluation metric allows us to compare systems:

"The basic system gets 81% accuracy, but after making some changes, the accuracy becomes 74%"



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#### **Evaluation**

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The evaluation metric allows us to compare systems:

"The basic system gets 81% accuracy, but after making some changes, the accuracy becomes 74%" — Why?



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Typically, comparison is more difficult:

"System A gets 45% precision and 80% recall; System B gets 95% precision and 10% recall"



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Typically, comparison is more difficult:

"System A gets 45% precision and 80% recall; System B gets 95% precision and 10% recall" — Which one should we use? (Also: why?)



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The answer depends on the problem (and the user)!



# Summary

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■ What is approximate string search?

- What are some common applications of approximate string search; why are they hard?
- What are some methods for finding an approximate match to a string? What do we need to generate them?
- How can we evaluate a typical approximate matching system?



# **Background Readings**

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# **Extension Readings**

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