Data analysis



Goals

- Be able to carry out simple text analysis tasks in the command line
- Be able to create a corpus in TXM
- Learn some of the basic functionalities of TXM for textual analysis:
 cooccurrences, specificity index, create partitions and subcorpora

Required software

The commands of this section are available in Unix-like operating systems. To enable them in Windows, you can easily install a Linux distribution if you have **Windows 10 version 2004 and higher (Build 19041 and higher) or Windows 11**

- Instructions (based on this post):
 - Control Panel > Programs > Turn Windows Features On Or Off. Enable the "Windows Subsystem for Linux" option in the list, and then click the "OK" button.
 - Alternatively, open PowerShell (<u>launching it as administrator</u>), and enter: wsl --install
 - After rebooting, go to Microsoft Store from the Start menu, and search for "Linux" and install Ubuntu.
 - Launch Ubuntu (from the start menu). The first time, you will be prompted to enter a UNIX username and password. Use this terminal to type the commands of the following slides

Count

Command:

wc [OPTION] [INPUT]

- It is used to find out number of lines, word count, byte and characters count in the files specified in the input arguments.
- By default, it displays four-columnar output.
 - 1 = number of lines
 - 2 = number of words
 - 3 = number of characters
 - \circ 4 = filename

Count

In the folder "datasets/plain-text/disco-19th" run the following commands:

- 1. wc *
- 2. wc * > ../count.csv (to print the results in a file)
- 3. wc -w * (to just count the number of words per file)
- 4. wc -1 * (to just count the number of lines per file)

Find strings

Command:

```
grep [OPTION] [PATTERN] [INPUT]
```

• Searches for a pattern in a file.

Find strings

In the folder "datasets/plain-text/disco-19th" run the following commands:

- 1. grep "amor[a-z]" *
- 2. grep "\sroj[ao]" *
- 3. grep "\Sroj[ao]" *
- 4. **grep -1** "**Sroj[ao]**" * (to just output the filename where the string occurs)
- 5. grep "^yo" *
- 6. grep -y "^yo" * (to ignore the case)
- 7. **grep** -v -y -n "s" **disco140n.txt** (-v displays the lines that do not match the pattern; -n adds the line number)

Get unique words

Commands:

uniq [OPTION] [INPUT[OUTPUT]]

It reports and filters out the repeated lines in a file.

sort [INPUT FILE]

It sorts a file, arranging the records in a particular order.

Get unique words

In the folder "datasets/plain-text/word-list" run the following commands:

- 1. sort mrs-dalloway_woolf.txt | uniq >
 mrs-dalloway_unique-words.txt
- 2. sort mrs-dalloway_woolf.txt | uniq -i >
 mrs-dalloway_unique-words.txt
- 3. sort mrs-dalloway_woolf.txt | uniq -ic >
 mrs-dalloway_unique-words-count.txt
- 4. sort mrs-dalloway_woolf.txt | uniq -id | uniq -i >
 mrs-dalloway_repeated-words.txt

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Required software

- TXM and the <u>TreeTagger extension</u>

Importing files in TXM

Task: Create a corpus in TXM using as source the <u>ELTEC-eng</u> corpus available in the folder "datasets/tei"

- Modify the file ELTeC-eng_metadata.tsv accordingly, so the metadata can be read in TXM
 - It must be a CSV file, with a comma as delimiter and named metadata.csv
 - The first column must be named "id" and contain the filenames (without extension)
 - The metadata.csv file must be saved in the same folder that contains the XML files (in the folder "exemplars" you have an example of a metadata.csv file with the ELTEC-eng data
- 2. Import the corpus with the option XML/w + CSV

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CQL

- Corpus Query Language = enables the search for complex grammatical or lexical patterns by accessing the properties annotated in each token
 - It supports regular expressions
- TXM offers help building CQL expressions, so you can carry out many queries without knowing the language.

CQL syntax

- Criteria for each token must appear between a pair of square brackets with the format:
 - o [attribute="value"]
- To search for a phrase, each token must appear in its own pair of square brackets:
 - [lemma="add"][word="salt"][word="to"][pos="NOUN"]
- Square brackets [] stand for 'any token'. Curly brackets { } are used for repetition of the preceding token.
 - [lemma="refill"] [] [lemma="teapot"]
 - [lemma="have"] []{2,4}[lemma="opinion"]

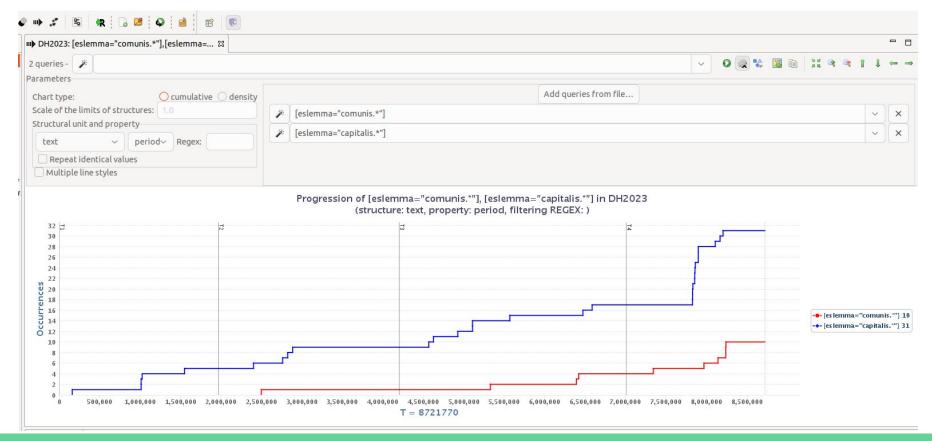
CQL syntax

- Combine different criteria with & (and) and | (or). E.g.:
 - o [lemma="power" & pos="VERB"]
 - [lemma="refill"] [] [lemma="kettle|teapot"]
- Negate with ! (not):
 - o [lemma="power" & pos!="V.*"] or [lemma="power" &
 !pos="V.*"]

Progression analysis

- Global visualisation of a development (corpus analysed as a continuum). It enables the quantitative evolution of a word/collocation, for instance, through.
- Results presented as a cumulative chart (need to interpret the slope of the curve)

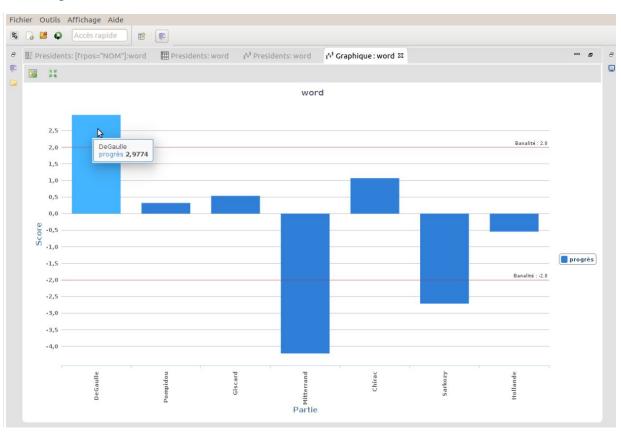
Progression analysis



Specificity index

- Based on the formula by Lafon (1984):
 - o Lafon, P. (1984). Dépouillements et statistiques en lexicométrie. Genève, Paris: Slatkine.
- It can be used to show the specificity of a cooccurrence or to characterise a corpus partition

Specificity index



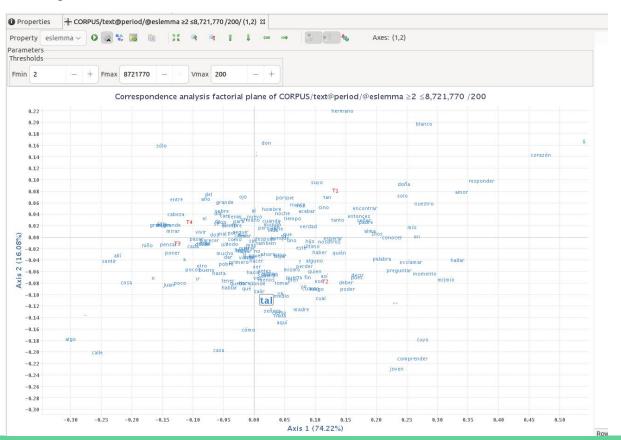
Task

- 1. Create a partition: one with the period T1 and another with the period T4
- 2. Calculate the specificities of PoS tags
- 3. Create a chart with the results

Factorial analyses

Cartographic representation of the distribution of a specific property across partitions

Factorial analyses



To learn more

- Very fast introduction to TXM key concepts in 90 minutes: <u>PDF</u>
- Import of Europresse articles in XML/w+CSV to TXM
- <u>List of manuals and resources in the TXM portal</u>