CHAINING SEARCH NOTEBOOK - QUICK START

The Chaining search notebook is an advanced digital environment for linguistic research. It supplies more than common tools like an graphic user interface for searching a lexicon or a corpus. The notebook provides the user with a set of functions allowing highly customizable search and data processing operations. One can write custom Python scripts to do multiple searches and process the results, draw charts, and of course 'chain' operations. That is: the results of the one search can be programmatically re-used in another search, in another source, in another way.

INSTALLATION

The installation procedure of the notebook is fully described in the README . md file on GitHub. Please refer to this file. Installation scripts are provided for both Linux/Mac and Windows.

LET'S START

Once the notebook is installed and launched as indicated in the README.md file, your browser will automatically open the following page:

💭 Jupyter	Quit Logout
Files Running Clusters Nbextensions	
Select items to perform actions on them.	Upload New - 2
0 - 1	Name Last Modified File size
C chaininglib	12 days ago
🔲 🗅 doc	21 days ago
env 🗅 env	20 days ago
Examples.ipynb	7 days ago 39.2 kB
🔲 릗 Sandbox.ipynb	Running 21 minutes ago 132 kB
install.bat	17 days ago 914 B
install.sh	18 days ago 1.09 kB
Makefile	21 days ago 598 B

As shown hereabove, the notebook consists of some folders and files. Only two files are relevant to you now:

- Examples.ipynb and
- Sandbox.ipynb

The first file, Examples.ipynb, is a notebook filled with case studies. These are examples showing how to perform search and data processing operations of several types and complexity.

Once these examples have inspired you, you'll be ready to go ahead with the second file, Sandbox.ipynb, which is an empty notebook, where you can try things out or build a complex script straight away.

The following sections will present both files and explain what you can do with those.



CASES STUDIES / EXAMPLES

The Examples.ipynb file can be accessed but just clicking on it. The file will open in a new tab called 'Examples' which should look like this:

💭 jupyte	er Examples (autosaved)
File Edit	View Insert Cell Kernel Widgets Help Trusted env O
₽ + ≫	
_	Examples Chaining search
Ť	Examples Chaining search
	This notebook contains a number of examples of chaining linguistic resources: corpora, lexica and
	treebanks. Try the examples, of copy the code and customize the examples in the <u>Sanobox</u> .
-	List of examples
	-
	Corpora
	<u>Corpus search</u>
	 Erequency of zeker+verb and vast+verb compared

Make sure it's working

First of all, when opening this file for the first time, we have to make sure that we are using the right kernel and that the kernel is trusted.

File	Edit	View	Insert	Cell	Kernel	Widgets	Help	Trusted	env O
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Here is how to do that:

- First: The name of the kernel in use is shown in the horizontal grayish top bar (see picture). On its right edge, it should show the word 'env'. If it does, you're ok. If it doesn't, click menu item 'Kernel' in the same bar and choose option 'Change kernel'; this will opens a pulldown menu where you must click the 'env' option. The kernel name on the right side of the bar must now read 'env' as well.
- Second: On the left side of the kernel name, there must be a small box saying 'Trusted'. If it does, we're done. But if it says 'Not trusted' instead: click on it, and when a confirmation window pops up, click on 'Trust'. Done!

OVERVIEW

Now we are ready to have a look at the examples. We have quite a list of those:

List of examples

Corpora

- <u>Corpus search</u>
- Frequency of zeker+verb and vast+verb compared
- <u>Train a POS tagger on an annotated corpus</u>
- Search in corpus and filter on metadata
- <u>Visualizing h-dropping</u>
- Generate lexicon from several corpora

Lexica

Lexicon search

Corpus + lexicon

- <u>Retrieve synonyms from DiaMaNT, look up in Gysseling</u>
- · Build a frequency list of the lemma of some corpus output
- Find occurences of attributive adjectives not ending with -e, even though they are preceeded by a definite article
- Look up inflected forms and spelling variants for a given lemma in a corpus
- <u>Corpus frequency list of lemmata from lexicon with given lemma</u>
- Build a frequency table of some corpus, based on lemmata of a given lexicon
- · Search corpus for wordforms of lemma not included in lexicon

Treebanks

- <u>Treebank search</u>
- Which objects of verb geven occur?

As the screenshot shows, the examples are ordered by category. You can visit an example straight by clicking on its name.

RUNNING A CASE STUDY

We've just seen the list of available case studies/examples. Now, clicking on the first example ('corpus search') will lead you to this screen section:

File Edit	View Insert Cell Kernel Widgets Help Trusted env O
🖹 🕇 🕅 🌾	A ↓ N Run ■ C > Markdown ▼
-	Corpus search
	Run the cell below to show the UI, and fill in your search query
	Choose one of the corpora:
	century letters from Dutch sailors
	gysseling: Corpus Gysseling, 13th century Dutch
	chn-extern: Externally accessible part of the Corpus Hedendaags Nederlands. Corpus of contamporant Dutch from the Dutch Antillee and Suriname, ratio and from neuroparate
	and websites.
	opus: OPUS corpus of Dutch subtitles
In []:	from chaininglib.ui.search import create_corpus_ui
	from chaininglib.ui.dfui import display_df, get_uploader
	# Create corpus UI, creates references to field contents
	<pre>corpusQueryField, corpusField = create_corpus_ui()</pre>

Each example is built quite the same way as above. The title of the example is followed by some explanation, an a cell with some Python code in it.

To run the code:

- Click into the cell
- Then go to the horizontal top bar and click on 'Run'

Once we've done that, some output should appear under the code cell. In this particular case, our output is a search interface, in which we can enter a search query and choose a corpus to search:

In [1]:	<pre>from chaininglib.ui.search import create_corpus_ui from chaininglib.ui.dfui import display_df, get_uploader</pre>						
	<pre># Create corpus UI, creates references to field contents corpusQueryField, corpusField = create_corpus_ui()</pre>						
	CQL query: [lemma="boek"]						
	Corpus: zeebrieven V						

To start the search, go to the next code cell.¹ This following piece of code reads your input out of the interface and start a search with that:



When this code cell is clicked upon and subsequently run, the search results will appear underneath as expected:

				ults	Resu
right conte	word 0	pos 0	lemma 0	left context	
en pampier en pennen	boeken	NOU	boek	heeft 4 gl 0 Aen	0
waet ghij uijt geft d	boeck	NOU	boek	lijeue man stelt alles te	1
en stondt en hij he	bouck	NOU	boek	alzoo hij niet op de	2
50 ditto ongsneeden ditto	boek.	NOU	boek	Schrijfpampier a 6 Sr t	3
bij berntrop laat verkoopen, a	boeken	NOU	boek	voorne missive, dat UEDs mijn	4
aan gaat, schouten altoos gepraferee	boeken	NOU	boek	verbeeld hebbe, dat wat geleerde	5
& Voorsiatiahaiid die priise	hoecken	NOU	boek	haiida accuratassa noada ordar dar	6

This is all you need to know to be able to explore the following example! For each example, click the code cell you want to run the code of, click on 'Run' in the horizontal top bar, and see the results.

Important remark

Note that in some rare cases, an example can only be run when the previous example was run before. But in those cases, the introduction text indicated that.

¹ A mistake commonly made in this particular case study is to start the search by clicking on 'Run' in the horizontal top bar again. But since the cell that contains the code for generating the search interface is still active (because that's the last element you clicked on or typed in), clicking on 'Run' again will just cause the interface to be rebuilt. So the search won't start at all. To be able to start the search, you need to go to the next code cell. This cell contains code that picks up the input from the interface and start the search with it.

YOU OWN SANDBOX

Now you're finished with the case studies, let's look at the Sandbox.ipynb file. That file is the place where we'll build our own scripts.

To access this file, we'll first have to go back to the browser tab called 'Home'. This tab shows the list of files or folders available in our notebook installation.

C Home		×	🗐 Examples	× +
ree				
💭 Jup	oyter			
Files	Running	Clusters	Nbextensions	
	 Image: Image: Ima	uons on them		
) chaininglib			
) env			
	Examples.ipyr	ıb		
	Sandbox.ipynl	b		
	install.bat			

Look up the Sandbox.ipynb file in the list and click on it. The file will open in a new tab and should look like this:

Jupyter Sandbox (unsaved changes)	4	Logout
File Edit View Insert Cell Kernel Widgets Help	Trusted	env O
E + ≫ @ T + ↓ H Run ■ C → Code		
 Chaining search: Sandbox 		
Use this notebook to combine linguistic resources yourself: corpora, lexica and	l treebanks.	
Use methods from our library chaininglib, described in the documentation		
 To get an idea of the possibilities and to copy code, go to the Examples not 	otebook.	
In []: import chaininglib		
In []:		

Just like in the Case studies screen, you have cells at your disposal. You can type your own code there, following what we've seen in the Case studies, and also the methods documentation. Both can be accessed again from this screen by clicking the corresponding link (see picture). Finally, click on the 'Run' button to execute your code.

Happy coding!