

How to use LexMapr

For people who have
little to no experience with command line

and

For macOS and Linux users only

and

For IFSAC users only

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Open up your terminal

macOs:

cmd + space to open spotlight search

type terminal

hit *return*

Linux:

ctrl + alt + t

Terminal tutorial

The terminal opens in your Home (also known as ~) directory

Type the following command into your terminal:

```
ls
```

This will print out all the sub-directories and files in your home directory

Terminal tutorial

You can change terminal directories with the `cd` command:

<code>cd Desktop</code>	Navigate to your Desktop subdirectory
<code>ls</code>	View the contents of your Desktop
<code>cd ..</code>	Navigate to the parent directory
<code>ls</code>	View the contents of your Home again

If you ever get lost, you can return Home with `cd ~`

Do you have Conda?

In your terminal, type:

```
conda -V
```

If you have Conda, the terminal will tell you your Conda version number

If you do not have Conda, the terminal will tell you the `conda` command was not found

Install Conda

Go to <https://docs.conda.io/en/latest/miniconda.html>

Install the appropriate **64-bit bash installer** for **Python 3.7**

In your terminal, navigate to the folder containing the installer

Assuming this was the Downloads folder:

```
cd ~
```

```
cd Downloads
```

Install Conda

In this folder, we run the installer through a terminal command

macOS:

```
bash Miniconda3-latest-MacOSX-x86_64.sh
```

Linux:

```
bash Miniconda3-latest-Linux-x86_64.sh
```

Follow the prompts to completion

Bioconda

Bioconda is a channel that allows users to easily install various bioinformatic packages, including LexMapr

Run the following terminal commands to set up Bioconda:

```
conda config --add channels default
```

```
conda config --add channels bioconda
```

```
conda config --add channels conda-forge
```

RUN THESE COMMANDS IN ORDER

Install LexMapr into a Conda environment

Enter the following commands from any directory:

```
conda create -n LexMapr lexmapr
```

```
conda activate LexMapr
```

```
python -m nltk.downloader all
```

Follow the prompts to completion

This creates a Conda environment called “LexMapr”, that contains a Bioconda installation of LexMapr

Running LexMapr

Whenever you want to run LexMapr, all you have to do is:

Start the terminal

Activate your LexMapr Conda environment:

```
conda activate LexMapr
```

Run LexMapr commands

```
lexmapr {path_to_output_file} -p ifsac
```

LexMapr input files

LexMapr takes input csv files with the following format:

SampleId,Sample

1,Chicken Breast

2,Baked Potato

3,Canned Corn

4,Frozen Yogurt

5,Apple Pie

LexMapr input files

Open up Microsoft Excel, Apple Numbers or LibreOffice Calc
Create a spreadsheet like so:

	A	B
1	SampleId	Sample
2	1	Chicken Breast
3	2	Baked Potato
4	3	Canned Corn
5	4	Frozen Yogurt
6	5	Apple Pie

Save as “small_simple.csv”

Running LexMapr

In the terminal, `cd` to the folder containing `small_simple.csv`

With the LexMapr Conda environment activated, try:

```
lexmapr small_simple.csv -p ifsac
```

Running LexMapr

A new file called `ifsac_output.tsv` was just created in the same directory as `small_simple.csv`

Open it up in Microsoft Excel, Apple Numbers or LibreOffice Calc:

	A	B	C	D	E
1	Sample_Id	Sample_Desc	Cleaned_Sample	Matched_Components	Third Party Classification
2	small_simple1	Chicken Breast	chicken breast	[chicken breast:foodon_00002703]	[chicken]
3	small_simple2	Baked Potato	baked potato	[potato (whole, baked):foodon_03302196]	[root/underground (tubers)]
4	small_simple3	Canned Corn	canned corn	[corn (canned):foodon_03302665]	[seeded vegetables (other)]
5	small_simple4	Frozen Yogurt	frozen yogurt	[frozen yogurt:foodon_03307445]	[dairy]
6	small_simple5	Apple Pie	apple pie	[apple pie:foodon_00002475]	[pome fruit]

Running LexMapr

You can specify your own output file name by running:

```
lexmapr {path_to_output_file} -p ifsac -o {path_to_output_file}
```

e.g.

```
lexmapr small_simple.csv -p ifsac -o small_simple_output.tsv
```

Now you results will be in `small_simple_output.tsv`, **not** `ifsac_output.tsv`

Keep LexMapr up-to-date

Keep LexMapr up-to-date by running the following command from any directory:

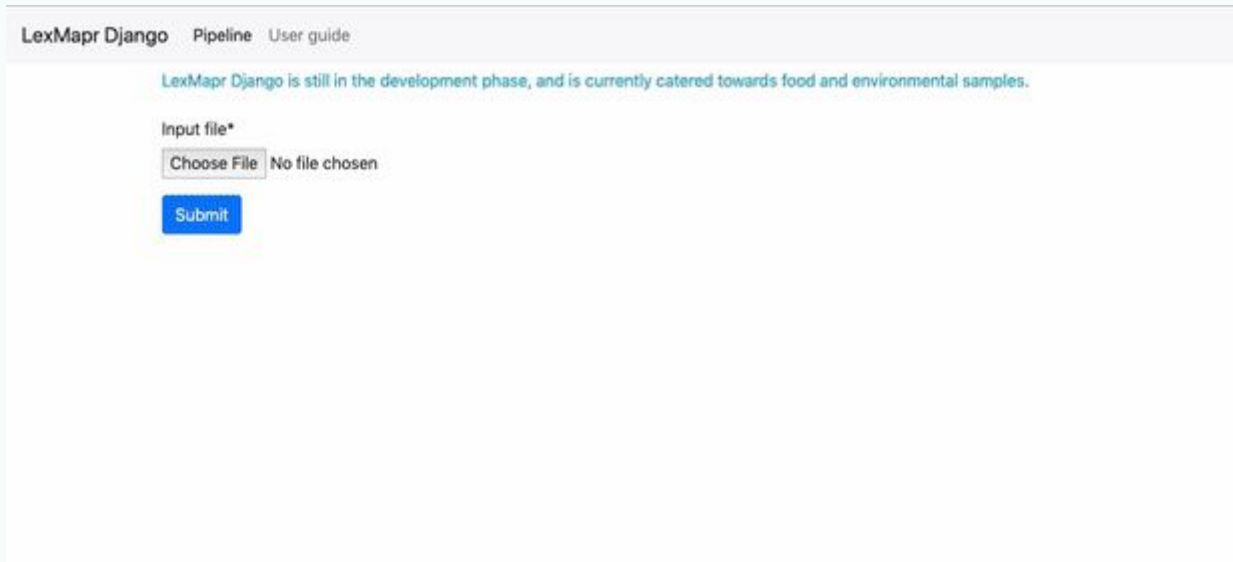
```
conda update lexmapr -n LexMapr
```

Follow the prompts to completion

This updates the lexmapr package installed inside your LexMapr conda environment

LexMapr Django

<https://watson.bccdc.med.ubc.ca/lexmapr/>



The screenshot shows the LexMapr Django web interface. At the top, there are navigation links: "LexMapr Django", "Pipeline", and "User guide". Below these, a teal message states: "LexMapr Django is still in the development phase, and is currently catered towards food and environmental samples." The main section is titled "Input file*" and contains a "Choose File" button next to the text "No file chosen". Below this is a blue "Submit" button.

Django-powered web interface, that allows you to run LexMapr from your browser