Efficient Literature Searches using Python

Blair Bilodeau May 30, 2020

University of Toronto & Vector Institute









Workshop Motivation



- Me trying to read all the new papers posted on arXiv

• Discuss the goal of focused literature searches v.s. reading new updates.

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- Practice!



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- Avoid keeping track of all new papers many will quickly become irrelevant to you.

Researchers in stats, math, bio, medicine, and other fields use these to post versions of papers before publication (as well as open source access after publication).

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Disadvantages

- No peer-review, so papers may be rougher.
- Easy to get lost in a sea of papers.

Preprint Server Search Options



	Q
e a term to search within all a	articles on this preprint server: e.g. stem cell
LIMIT RESULTS	
Date Posted	
	m —
ýpe a term to search within a	W articles on this preprint server; e.g. stem cell
	m
ype a term to search within a	W articles on this preprint server: e.g. stem cell
Include articles in bioRxiv and/or	bioRxiv × _
medRxiv:	UNIOTORY =
Include articles in	
subject area(s)	
	All Collections Animal Behavior and Cognition
Include articles of type:	Biochemistry
-77	Bioengineering
AUTHORS, KEYWOR	Bioinformatics Biophysics
earch for specific authors as	
Author	Cell Biology
	Clinical Trials
	m r r r r r r r r r r r r r r r r r r r
Salar a team to sensely within a	Developmental Biology
	W articles on Type a term to search within all articles
	W articles on Type a term to search within all articles
his preprint server: e.g. stern o	W articles on Type a term to search within all articles
his preprint server: e.g. stern o	W articles on Type a term to search within all articles
his preprint server: e.g. stem o	Warticles on Type a term to search within all articles ell this preprint server; e.g. stem cell words
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his preprint server: e.g. stem of FICIE Spe a term to search within a erver: e.g. stem cell	Warticles on Type a term to search within all articles ell this preprint server; e.g. stem cell words
his preprint server: e.g. stem of FICIE Spe a term to search within a erver: e.g. stem cell	If a stoke on . Type a term is seenth within oil arruche oil
his preprint server; e.g., stem of title Spe a term to search within a rever; e.g. stem cell Abstract or Title	of a residue on . Type a term is a server, within of a route of the properties according to the properties according to the properties according to the properties on this properties. words words
type a term to search within a his preprint server; e.g. stem o TICLE Type a term to search within a prever; e.g. stem cell Abstract or Title	of a residue on . Type a term is server to attitud of unusual and the propriet server of stem collection of the propriet server of stem collection of unusual and
his preprint server; e.g. stem of the server; e.g. stem of the server; e.g. stem cell serve	of a residue on . Type a term is a server, within of a route of the properties according to the properties according to the properties according to the properties on this properties. words words
his preprint server; e.g. stem of FITTLE Speed term to search within a rever; e.g. stem cell Abstract or Title Speed term to search within a erver; e.g. stem cell	of a recise on Type a term is a server by within all articles of this propriet server or g, stem coll who propriet a very 0 all phrase words If a recise on this propriet any 0 all phrase any 0 all phrase any 0 all phrase
his preprint server; e.g. stem of FITTLE Speed term to search within a rever; e.g. stem cell Abstract or Title Speed term to search within a erver; e.g. stem cell	If a relative on Tripe a form to secrets within of a route of the property according to the property according to the property according to the property of a secret on this property or any all phrase words on this property any all phrase or this property any all phrase or Tricle
his preprint server; e.g., stem of title Spe a term to search within a rever; e.g. stem cell Abstract or Title	of a recise on Type a term is a server by within all articles of this propriet server or g, stem coll who propriet a very 0 all phrase words If a recise on this propriet any 0 all phrase any 0 all phrase any 0 all phrase

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Biorxiv Options

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Biorxiv Options

No known options to me, besides this project with a broken link.
 (https://github.com/gokceneraslan/biorxiv-sanity-preserver)

Goals

- High flexibility for keyword searching.
- Easy to run and parse output everyday.
- Modular to allow for additional features to be added.

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Access the Scripts

https://github.com/blairbilodeau/arxiv-biorxiv-search

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If you don't see the following, you have to install.

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[blairbilodeau@Blairs-MacBook-Pro ~ % python3
Python 3.7.5 (v3.7.5:5c02a39a0b, Oct 14 2019, 18:49:57)
[Clang 6.0 (clang-600.0.57)] on darwin
Type "help", "copyright", "credits" or "license" for more information.
>>> |
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```

If you do see that, great! You're now in a python environment. Either spend some time in there (try typing print('hello world!')) or type exit() to leave. Take a break for the next slide.

Option 1: Directly Download Python

Go to https://www.python.org/downloads/ and download Python 3. (The actual version doesn't matter as long as it's Python 3.x.x)

Option 2: Use Anaconda

Download from https://www.anaconda.com/products/individual. (Preferable if you aren't familiar with working on the command line)

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Common Troubleshooting Tips

- Make sure the default python path is python3 if you have both installed
- On Windows, add python to your path environment
 - Computer: Properties: Advanced System Settings: Environment Variables: Path: add ";C:\Python36" (or whichever version) to the end

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Installing Packages

We will use pip, which is automatically included with installations.

To install a package named name:

open up terminal or command prompt and type pip install name.

On windows, you may need to type something like

 ${\tt C:\Python36\Scripts\pip\ install\ name\ or\ }$

C:\Python36\Scripts\pip.exe install name

For example, to install the package pandas,

```
|blairbilodeau@Blairs-MacBook-Pro ~ % pip install pandas
Collecting pandas
 Downloading pandas-1.0.3-cp37-cp37m-macosx_10_9_x86_64.whl (10.0 MB)
                                   10.0 MB 8.5 MB/s
Requirement already satisfied: python-dateutil>=2.6.1 in /Library/Frameworks/Pyt
hon.framework/Versions/3.7/lib/python3.7/site-packages (from pandas) (2.8.0)
Requirement already satisfied: numpy>=1.13.3 in /Library/Frameworks/Python.frame
work/Versions/3.7/lib/python3.7/site-packages (from pandas) (1.17.2)
Requirement already satisfied: pvtz>=2017.2 in /Library/Frameworks/Pvthon.framew
ork/Versions/3.7/lib/python3.7/site-packages (from pandas) (2019.3)
Requirement already satisfied: six>=1.5 in /Library/Frameworks/Python.framework,
Versions/3.7/lib/python3.7/site-packages (from python-dateutil>=2.6.1->pandas)
1.12.0)
Installing collected packages: pandas
Successfully installed pandas-1.0.3
blairbilodeau@Blairs-MacBook-Pro ~ %
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```
| blairbilodeau@8lairs-MacBook-Pro ~ % pip install pandas | Collecting pandas | Collecting pandas | Downloading pandas-1.0.3-cp37m-macosx_10_9_x86_64.whl (10.0 MB) | | 10.0 MB 8.5 MB/s | NB 8.5 MB/s
```

Extra packages needed for this script...

- pandas (data structure tools)
- requests (handling opening websites)
- beautifulsoup4 (parsing HTML)

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Script Idea

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- Repeat this for each subject;
- Display the titles and abstracts selected (with other info if desired), with optional exporting of information to csv file and downloading of full pdfs.

```
#######
## Main function
## Parameters:
# sdate

    datetime object

                                       - starting date of search period
                                       - end date of search period
# fdate
# kwd_rea
                                       - keywords that are required to be in the title or abstract
                                       - keywords that must not be in the title or abstract
             - list of lists of str - keywords of which at least one must be included
             - list of str
                                       - authors which are required
# athr_exc
             - list of str
                                       - authors to exclude
# athr_one
             - list of lists of str - authors of which one must be included
             - list of str
                                       - subject options are:
                                           astro-ph, cond-mat, ar-ac, hep-ex, hep-lat, hep-ph, hep-th, math-ph, nlin, nucl-ex, nucl-th,
                                           physics, quant-ph, math, cs, a-bio, a-fin, stat
                                       - maximum number of results to return
# max_time
                                       - maximum amount of seconds to be spent searching
# cols
                                           column options are:
# export
                                       - folder location to dump results list in csv
# exportfile - str
                                       - by default the file name will be today's date, but you can override this with exportfile
# download
                                       - folder location to dump returned pdfs into
                                           (files named using year_lastname format)
```

- kwd_req, kwd_exc, kwd_one are the main parameters that allow for custom searching of papers
- All of these are optional if you don't pass any arguments you will get the first 50 papers from cs for the month

arxiv_search_function.py Demonstration

I will now show an example of running through the code in a Jupyter notebook.

 $\verb|biorxiv_search_function.py| \textbf{Parameters}|$

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```
****
## Main function
# sdate
              - datetime object
                                        - starting date of search period
# fdate
               - datetime object
                                        - end date of search period
# kwd
                                        - keywords to search for in title and abstract
# kwd_tvpe
             - 'all' or 'anv'
                                        - whether all keywords are required or just one of them
# athr
# subjects
                                        - subject options are:
                                          (Note capitalization and spacing are important for subject)
                                            Animal Behaviour and Coanition, Biochemistry, Bioengineering, Bioinformatics,
                                            Ecology, Epidemiology, Evolutionary Biology, Genetics, Genomics, Immunology,
                                            Micropiglogy, Molecular Biology, Neuroscience, Paleontology, Pathology,
                                            Pharmacology and Toxicology, Physiology, Plant Biology, Scientific Communication and Education
                                            Synthetic Biology, Systems Biology, Zoology
                                        - maximum amount of seconds to be spent searching
# cols

    bidrxiv fields to extract

                                            column options are:
                                            title, authors, date, url
# export
                                        - folder location to dump results list in csv
# exportfile
                                        - by default the file name will be today's date, but you can override this with exportfile
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```

Timesaving Workflow

Running the Script

- Create a separate python file to call the functions with parameters you desire, and run that from command line every day.
- See the file search_examples.py in my Github.

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Automating the Script

- Mac: used launchd
 - Create a shell script to run the python file you want (search_examples.sh)
 - Place the file file.name.plist in /Library/LaunchDaemons with names changed (currently runs once every 24 hours, can be changed).
 - In command line, type cd Library/LaunchDaemons and then sudo launchctl load file.name.plist.
- Windows: use Task Scheduler
 - Create a batch file to run the python file you want.
 - Follow the instructions in Task Scheduler after clicking Create Basic Task.