

Introduction to NCBI EDirect Software

Learn how to programmatically search and compile PubMed and related data in a Unix Shell

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Disclaimer: Workshop is not affiliated with NCBI/NLM/NIH

Outline

1. Overview of NCBI EDirect
2. Brief Introduction to Unix Shell
3. Searching PubMed and related databases with EDirect
4. Retrieving PubMed XML and extracting information
5. Creating scripts for repeated tasks and workflows

Slides and Further Reading

These slides (.pdf) are available (MIT license):

<https://github.com/vfscalfani/EDirectChemInfo>

Go to **Workshops** folder.

Key References and Further Reading:

1. [Software Carpentry: The Unix Shell](#)
2. Official NCBI Manual for EDirect - [Entrez Direct: E-Utilities on the Unix Command Line](#)
3. NLM EDirect for PubMed Recordings and Materials - [EDirect for PubMed](#)
4. NLM EDirect Documentation on [xtract](#)
5. [NCBI EDirect Cookbook](#)
6. Our PubMed/PubChem EDirect Cookbook, [EDirectChemInfo](#) and [Unix Introduction](#)

Appropriate EDirect and NCBI Data Usage Notes

Read the NCBI Website and Data Usage Policies and Disclaimers:

<https://www.ncbi.nlm.nih.gov/home/about/policies/>

See information about abstract copyright in PubMed:

<https://www.nlm.nih.gov/databases/download.html>

And PubMed Central Copyright Notice:

<https://www.ncbi.nlm.nih.gov/pmc/about/copyright/>

If you have questions about copyright and fair-use for your particular use-case, please contact The University of Alabama Libraries: <https://ask.lib.ua.edu/>

What is EDirect [1]?

- Free command-line program from National Center for Biotechnology Information (NCBI) that allows (E-utilities) programmatic access to NCBI databases such as PubMed, PubChem, Gene, Taxonomy, etc. directly within a Unix terminal window.
- Can be installed on Unix, Unix-like (e.g., GNU/Linux) distributions, Mac OS, and Windows with Cygwin Unix-emulation.

Example EDirect Use

```
$ esearch -email name@xx.edu -db pubmed -query "\"ionic liquids\"[MESH] AND imidazolium" | \
> efetch -format xml | \
> xtract -pattern PubmedArticle -element MedlineCitation/PMID -first Author/LastName \
> Author/Initials ISOAbbreviation PubDate/Year Volume Issue MedlinePgn \
> -block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
33396149          Hu          LX          Ecotoxicol Environ Saf          2021    208    111629
https://doi.org/10.1016%2Fj.ecoenv.2020.111629
33346267          Kaur          M          Phys Chem Chem Phys          2021    23    1    320-328 https://doi.org/10.1039%2Fd0cp04513f
33253998          Tashakkori P          J Chromatogr A 2021    1635    461741    https://doi.org/10.1016%2Fj.chroma.2020.461741
...
...
```

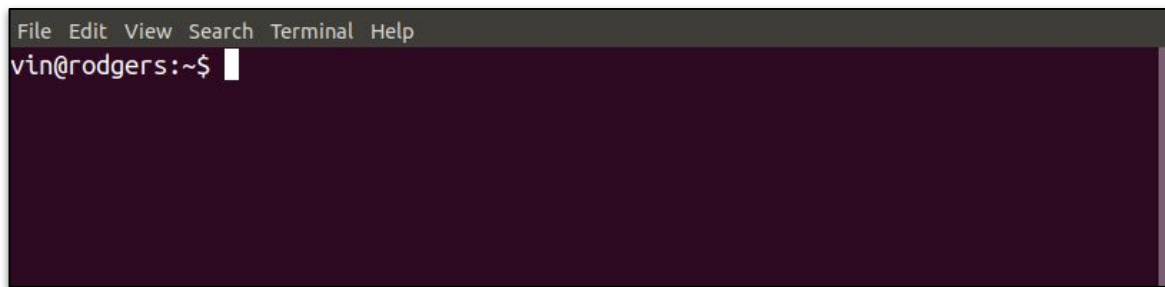
[1] <https://www.ncbi.nlm.nih.gov/books/NBK179288/>

Why Would I Use EDirect?

1. For accessing NCBI data programmatically, EDirect has a lower learning curve than trying to write your own scripts in a programming language. EDirect constructs the programmatic web URLs for you and includes programs to help you format and process the data into tabular formats. *Can also combine/process data with other Unix utilities.*
2. You want to compile bibliographic data or molecular/biological datasets.
3. You have many search queries (e.g., PubMed) to perform or need to repeat the search frequently. Easy to precisely record your database searches and analysis, which may be useful for systematic reviews.
4. You can quickly answer interesting and specific questions like “Who are the most common authors studying peanut allergies?” or “What is the most common journal indexed in PubMed for research on synthetic chemistry “total synthesis”?”
5. Searching NCBI databases in a Unix terminal is a lot of fun.

Unix Terminal

A Unix terminal is a text input/output environment [1]:



From the terminal input, a shell then interprets the commands (i.e., a command line interpreter).

Most Unix-like operating systems such as GNU/Linux distributions (e.g., Ubuntu) are using the [GNU Bash Shell](#).

[1] Unix Stack Exchange Thread: [What is the exact difference between a 'terminal', a 'shell', a 'tty' and a 'console'?](#)

Unix Programs and Utilities

To run a Unix program, you generally type the name of the program, followed by (optional) -arguments.

Type **-help** or **--help** after the program name or **man** (for manual) before the program name to see specific usage. Example with GNU utilities `cut`:

```
$ cut --help
```

```
Usage: cut OPTION... [FILE]...
```

```
Print selected parts of lines from each FILE to standard output.
```

```
With no FILE, or when FILE is -, read standard input.
```

```
Mandatory arguments to long options are mandatory for short options too.
```

```
-b, --bytes=LIST      select only these bytes
-c, --characters=LIST  select only these characters
-d, --delimiter=DELIM use DELIM instead of TAB for field delimiter
-f, --fields=LIST      select only these fields; also print any line
                       that contains no delimiter character, unless
                       the -s option is specified
```

```
...
```

```
$ man cut
```

```
(outputs manual page for cut, more detailed description, not shown)
```


Unix Shell Pipelines, Redirect, and Loops [1]

With the Unix shell, we can use pipelines to create sequences of commands. Each command output is piped into the next command:

```
$ command1 | command2 | command3
```

We can redirect our output from a command or sequence of commands to a file:

```
$ command1 > myfile1.txt
```

```
$ command1 | command2 | command3 > myfile3.txt
```

Unix shell is also a programming language, and, for example, we can create loops to repeat tasks:

```
$ for item in list_of_items  
> do  
>     something_using $item  
> done
```

[1] See the [Software Carpentry Unix Shell](#) and [Bash Reference Manual](#)

EDirect Unix Programs [1]

EDirect contains several individual programs. We will review the following today:

1. **einfo** - prints fields and links indexed in each database
2. **esearch** - performs an NCBI Entrez database search based on a specified database and query
3. **efetch** - downloads the esearch query results in a specified format such as XML
4. **xtract** - extracts selected data values from XML
5. **elink** - finds associated records within a specified database
6. **efilter** - limits results (e.g., by date, information type, etc.)

Typical use-case is to connect these programs with unix pipelines:

```
$ esearch | efetch | xtract
```

```
$ esearch | elink | efilter | efetch | xtract
```

```
$ esearch | elink | efilter | efetch | xtract | sort | uniq -c
```

```
$ esearch | elink | efilter | efetch | xtract > myfile.txt
```

[1] See the official manual for other EDirect programs and examples: <https://www.ncbi.nlm.nih.gov/books/NBK179288/>

EDirect program syntax and Usage Notes

EDirect programs all have similar syntax:

```
eprogram -argument input
```

```
eprogram -email name@xx.edu -argument input
```

All of the EDirect programs accept your email as an option too, this is a really good idea to add so that if you are accidentally causing server issues or violating their usage policies, NCBI can contact you.

See earlier slide 4 entitled, "Appropriate EDirect and NCBI Data Usage Notes."

einfo

einfo - prints fields and links indexed in each database

```
einfo -help
```

```
einfo -dbs
```

```
einfo -db pubmed -fields
```

```
einfo -db pubmed -links
```

EDirect manual: <https://www.ncbi.nlm.nih.gov/books/NBK179288/>

esearch

esearch - performs an NCBI Entrez database search based on a specified database and query

```
esearch -help
```

```
esearch -db pubmed -query "17630804"[UID]
```

```
esearch -db pubmed -query "imidazolium AND bacteria"
```

As queries become more complex, use the `-debug` flag to check the query translation:

```
esearch -email vfscalfani@ua.edu -db pubmed -query "hydrogel-based drug delivery" -debug
```

```
nquire -url https://eutils.ncbi.nlm.nih.gov/entrez/eutils/ esearch.fcgi -retmax 0 \  
-usehistory y -db pubmed -term "hydrogel-based drug delivery"
```

esearch

esearch - performs an NCBI Entrez database search based on a specified database and query

Escape `,` `\`, internal quotes and use parentheses for complex searches:

```
esearch -db pubmed -query "\"Artificial Intelligence\"[MESH] AND \"drug discovery\"[ALL]"
```

```
esearch -db pubmed -query "(university of alabama[AFFL]) NOT (birmingham[AFFL] OR \
huntsville[AFFL])"
```

Again, try the `-debug` flag for testing, and it is also helpful to build queries online with the PubMed Advanced Search Builder: <https://pubmed.ncbi.nlm.nih.gov/advanced/>, though **e-utilities based searches may be different than the web based PubMed.**

efetch

efetch - downloads the esearch query results in a specified format such as XML

```
efetch -help
```

```
esearch -db pubmed -query "17630804"[PMID] | \  
efetch -format abstract
```

```
esearch -db pubmed -query "17630804"[PMID] | \  
efetch -format xml
```

```
esearch -db pubmed -query "\"Artificial Intelligence\"[MESH] AND \"drug discovery\"[ALL]" | \  
efetch -format xml
```

EDirect manual: <https://www.ncbi.nlm.nih.gov/books/NBK179288/>

xtract

xtract - extracts selected data values from XML

```
xtract -help
```

Very powerful tool, we will look at some basics today.

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml
```

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml | \
```

xtract -outline

Let's take a closer look at the PubMed XML

EDirect manual: <https://www.ncbi.nlm.nih.gov/books/NBK179288/>

PubMed XML example

```
1 <?xml version="1.0" encoding="UTF-8" ?>
2 <!DOCTYPE PubMedArticleSet PUBLIC "-//NLM//DTD PubMedArticle, 1st January 2019/
3 <PubMedArticleSet>
4   <PubMedArticle>
5     <MedlineCitation Status="MEDLINE" Owner="NLM">
6       <PMID Version="1">17630804</PMID>
7       <DateCompleted>
8         <Year>2007</Year>
9         <Month>10</Month>
10        <Day>25</Day>
11      </DateCompleted>
12      <DateRevised>
13        <Year>2007</Year>
14        <Month>08</Month>
15        <Day>10</Day>
16      </DateRevised>
17      <Article PubModel="Print-Electronic">
18        <Journal>
19          <ISSN IssnType="Print">0022-3263</ISSN>
20          <JournalIssue CitedMedium="Print">
21            <Volume>72</Volume>
22            <Issue>17</Issue>
23            <PubDate>
24              <Year>2007</Year>
25              <Month>Aug</Month>
26              <Day>17</Day>
27            </PubDate>
28          </JournalIssue>
29          <Title>The Journal of organic chemistry</Title>
30          <ISOAbbreviation>J Org Chem</ISOAbbreviation>
31        </Journal>
32        <ArticleTitle>Total synthesis and absolute configuration determination
33        <Pagination>
34          <MedlinePgn>6621-3</MedlinePgn>
35        </Pagination>
```

```
1 PubMedArticle
2   MedlineCitation
3     PMID
4     DateCompleted
5       Year
6       Month
7       Day
8     DateRevised
9       Year
10      Month
11      Day
12   Article
13     Journal
14       ISSN
15     JournalIssue
16       Volume
17       Issue
18     PubDate
19       Year
20       Month
21       Day
22   Title
23   ISOAbbreviation
24   ArticleTitle
25   Pagination
26     MedlinePgn
```

Xtract [1]

Basic usage today:

\$ xtract **-pattern** A **-element** B C...

Key concepts:

1. pattern defines new rows (e.g., PubMedArticle, Author)
2. element defines new columns (e.g., ArticleTitle, Volume, Issue...)
3. Attributes of XML elements (e.g., <PMID **Version**="1">17630804</PMID>) can be selected with @:
 - a. PMID@Version
4. In cases where elements have the same name (e.g., Year), use a / to define your selection as Parent/Child hierarchy
 - a. PubDate/Year versus DateRevised/Year

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>

xtract Examples

extract author names as 1 author per line [1]

```
esearch -db pubmed -query "17630804"[PMID] | \  
efetch -format xml | \  
xtract -pattern Author -element Author/LastName Author/Initials
```

extract author names as 1 article per line [1]

```
esearch -db pubmed -query "17630804"[PMID] | \  
efetch -format xml | \  
xtract -pattern PubmedArticle -element Author/LastName Author/Initials
```

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>

xtract Examples

Extract PMID and other bibliographic information [1]

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -element MedlineCitation/PMID ArticleTitle \
ISOAbbreviation PubDate/Year Volume Issue MedlinePgn
```

Add in author names [1]

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -element MedlineCitation/PMID Author/LastName \
Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue MedlinePgn
```

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>

xtract Examples

Reformat author names using the xtract -block argument [1]

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -element MedlineCitation/PMID ArticleTitle ISOAbbreviation \
PubDate/Year Volume Issue MedlinePgn \
-block Author -element LastName Initials
```

extract only first author names using the xtract -first argument [2]

```
esearch -db pubmed -query "17630804"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -element MedlineCitation/PMID -first Author/LastName \
Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue MedlinePgn
```

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>; [2] <https://github.com/vfscalfani/EDirectChemInfo>

xtract Examples

Good idea to use the xtract default field (-def) value to handle missing fields [1]

```
esearch -db pubmed -query "25818947"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first \
Author/LastName Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue \
MedlinePgn
```

Add in the DOI with the -block and conditional -if [1]

```
esearch -db pubmed -query "25818947"[PMID] | \
efetch -format xml | \
xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first \
Author/LastName Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue
MedlinePgn \
-block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
```

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>

xtract Examples

the same xtract commands can work for queries that return multiple articles [1,2]

```
esearch -db pubmed -query "Anthraquinones/chemical synthesis" [MESH] | \  
efetch -format xml | \  
xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first Author/LastName \  
Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue MedlinePgn \  
-block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
```

[1] <https://dataguide.nlm.nih.gov/edirect/xtract.html>; [2] <https://github.com/vfscalfani/EDirectChemInfo>

elink

elink - finds associated records within a specified database

Citation information from [NIH Open Citation Collection](#)

```
elink -help
```

```
elink -cited      (references to this article)
elink -cites      (article reference list)
elink -related    (neighbors in same database)
```

Get article citations for PMID 31254167

```
esearch -db pubmed -query "31254167[PMID]" | \
elink -cited | \
efetch -format xml | \
xtract -pattern PubmedArticle -element MedlineCitation/PMID -first Author/LastName \
Author/Initials ISOAbbreviation PubDate/Year Volume Issue MedlinePgn \
-block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
```


elink Example

elink - finds associated records within a specified database

Can also specify another database:

```
elink -target new_database -name related-link
```

Recall: `einfo -db pubmed -links`

Get related PubChem compounds from a PubMed search [1]

```
esearch -db pubmed -query "Anthraquinones/chemical synthesis"[MESH] | \  
elink -target pccompound -name pubmed_pccompound | \  
efetch -format docsum | \  
xtract -pattern DocumentSummary -element IsomericSmiles CID InChIKey
```

[1] <https://github.com/vfscalfani/EDirectChemInfo>

efilter

efilter - limits results (e.g., by date, information type, etc.)

```
efilter -help
```

Basic example use:

```
efilter -query  
efilter -pub review  
efilter -mindate 2017
```

Sometimes these can be incorporated directly into esearch.

efilter examples

limit PubMed results to review articles only

```
esearch -db pubmed -query "Anthraquinones/chemical synthesis"[MESH] | \  
efilter -pub review | \  
efetch -format xml | \  
xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first \  
Author/LastName Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue \  
MedlinePgn -block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
```

limit PubMed results from a linked PubChem search to a specific Journal [1]

```
esearch -db pccompound -query 174076[uid] | \  
elink -target pubmed -name pccompound_pubmed | \  
efilter -query "Phys Chem Chem Phys"[JOUR] | \  
efetch -format xml | \  
xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first \  
Author/LastName Author/Initials ArticleTitle ISOAbbreviation PubDate/Year Volume Issue \  
MedlinePgn -block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId
```

[1] <https://github.com/vfscalfani/EDirectChemInfo>

Creating a For Loop for multiple Queries [1,2]

Let's say I have a list of PMIDs and want bibliographic information for each one:

```
for refs in \  
    "20426451" \  
    "21982300" \  
    "21948594" \  
    "12653513" \  
    "11259830" \  
    "10592235" \  
    "16796559" \  
    "27899562" \  
    "26400175" \  
    "8709122"  
  
do  
    esearch -db pubmed -query "$refs[PMID]" |  
    efetch -format xml |  
    xtract -pattern PubmedArticle -def "N/A" -element MedlineCitation/PMID -first Author/LastName \  
    Author/Initials ISOAbbreviation PubDate/Year Volume Issue MedlinePgn \  
    -block ArticleId -if ArticleId@IdType -equals doi -doi ArticleId  
    sleep 1  
  
done
```

Creating a For Loop for multiple Queries [1,2]

Or maybe we want the number of cited references for each PMID:

```
for refs in \  
    "20426451" \  
    "21982300" \  
    "21948594" \  
    "12653513" \  
    "11259830" \  
    "10592235" \  
    "16796559" \  
    "27899562" \  
    "26400175" \  
    "8709122"  
do  
    esearch -db pubmed -query "$refs[PMID]" |  
    elink -cited |  
    xtract -pattern ENTREZ_DIRECT -lbl "$refs" -element Count  
    sleep 1  
done
```

Answering Specific Questions

most common UA chemistry authors indexed in PubMed [1]:

```
esearch -db pubmed -query "(university of alabama[AFFL] AND tuscaloosa[AFFL])" | \
efetch -format xml | \
xtract -pattern Author -if Affiliation -contains chemistry -and Affiliation -contains \
tuscaloosa -element LastName Initials | \
sort-uniq-count-rank
```

[1] N.B. affiliation query and xtract pattern is not perfect, see more here:

https://github.com/vfscalfani/EDirectChemInfo/blob/master/05_EDirect_PubMed_Recipes.md

[2] sort-uniq-count-rank: <https://dataguide.nlm.nih.gov/edirect/sort-uniq-count-rank.html>

Answering Specific Questions

most Frequent Journals for a PubMed Query [1]

```
esearch -db pubmed -query "\"Artificial Intelligence\"[MESH] AND \"drug discovery\"[ALL]" | \
efetch -format xml | \
xtract -pattern PubmedArticle -element ISOAbbreviation | \
sort-uniq-count-rank
```

[1] <https://github.com/vfscalfani/EDirectChemInfo>

Answering Specific Questions

how many records are being added to PubMed by create date each month? [1]

```
for date in \  
    "2020/01" \  
    "2020/02" \  
    "2020/03" \  
    "2020/04" \  
    "2020/05" \  
    "2020/06"  
do  
    esearch -db pubmed -query "$date[CRDT]" |  
    xtract -pattern ENTREZ_DIRECT -lbl "$date" -element Count  
    sleep 1  
done
```

[1] <https://github.com/vfscalfani/EDirectChemInfo>

Answering Specific Questions

number of records for a PubMed query that are available in PubMed Central [1]:

```
esearch -db pubmed -query "J Chem Inf Model[JOUR]" | \  
elink -target pmc -name pubmed_pmc | \  
efetch -format docsum | \  
xtract -pattern DocumentSummary -element PubDate | \  
cut -d " " -f 1 | \  
sort-uniq-count-rank | \  
sort -k2,2
```

[1] <https://github.com/vfscalfani/EDirectChemInfo>

Answering Specific Questions

most frequent article title words from a PubMed query

```
esearch -db pubmed -query "J Cheminform[JOUR]" | \  
efetch -format xml | \  
xtract -pattern PubmedArticle -element ArticleTitle | \  
tr '\n' ' ' | \  
word-at-a-time | \  
sort-uniq-count-rank > titlewords.txt
```

And many more possibilities, use your imagination and look at linked resources on the next slide.

Thanks!

Key References and Further Reading:

1. [Software Carpentry: The Unix Shell](#)
2. Official NCBI Manual for EDirect - [Entrez Direct: E-Utilities on the Unix Command Line](#)
3. NLM EDirect for PubMed Recordings and Materials - [EDirect for PubMed](#)
4. NLM EDirect Documentation on [xtract](#)
5. [NCBI EDirect Cookbook](#)
6. Our PubMed/PubChem EDirect Cookbook, [EDirectChemInfo](#) and [Unix Introduction](#)

Need help?

Get in touch!

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