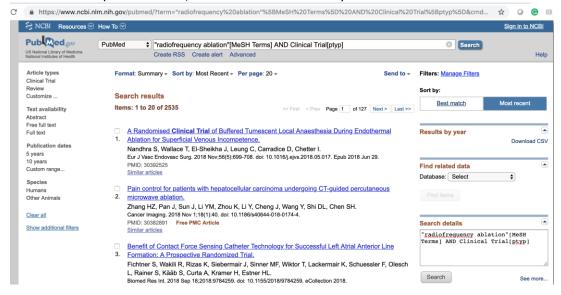
How to use pubmed API:

1. Brief guide: this is how to search in pubmed for common user:

For example, we want to search clinical trial of radiofrequency ablation.



Search Result (2535 results, first PMID is 30392525):



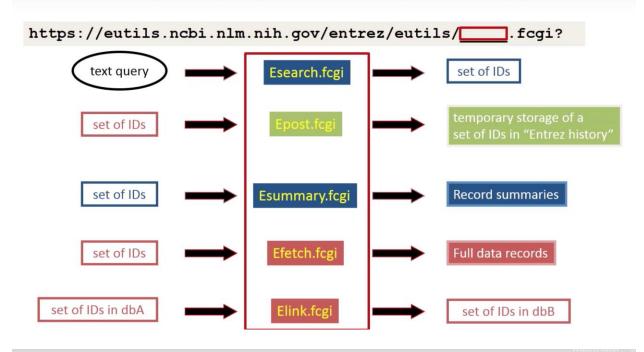
Search details of the right lower corner: radiofrequency ablation"[MeSH Terms] AND Clinical Trial[ptyp]

2. How to use pubmed API to search

according to the pubmed API, link:

https://www.ncbi.nlm.nih.gov/books/NBK25498/#chapter3.Application 4 Finding unique se it is the NCBI site-wide API: Entrez Programming Utilities(EUtils), use a set of structured URL-based calls to access the NCBI Entrez search system and databases.

VERBS (do what?) AND NOUN (with what?)



EUtils API Examples

Getting ESummary information for Gene with a Gene ID of 3269 (Human Histamine Receptor H1)

https://eutils.ncbi.nlm.nih.gov/entrez/eutils/esummary.fcgi?db=gene&id=3269

BASE URL CALL Perform this action on this database with this term

(return the information in this format)

Getting ESummary information for Gene with a Gene ID of 3269 (Human Histamine Receptor H1) https://eutils.ncbi.nlm.nih.gov/entrez/eutils/esummary.fcgi?db=gene&id=3269

Linking to Find PubMed Record IDs (PMIDs) for Gene ID 3269

/elink.fcgi?dbfrom=gene&id=3269&dbto=pubmed

Getting ESummary or Efetch information for PMID 24236486 (which was linked to Gene ID 3269)

<u>/esummary.fcgi?db=pubmed&id=24236486</u> (retype= xml is the default)

For our project, we can search Pubmed with a term(i.e: radiofrequency ablation, pain management, patient's diagnoses, clinical trial) and retrieve PMIDs(unique ID for each paper). We can also get summary for each paper using PMID.

there is an example how to search a term in pubmed from https://www.ncbi.nlm.nih.gov/books/NBK25497/

ESearch – ESummary/EFetch

Input: Entrez text query

ESummary Output: XML Document Summaries

EFetch Output: Formatted data records (e.g. abstracts, FASTA)

```
use LWP::Simple;
# Download PubMed records that are indexed in MeSH for both asthma and
# leukotrienes and were also published in 2009.
$db = 'pubmed';
$query = 'asthma[mesh]+AND+leukotrienes[mesh]+AND+2009[pdat]';
#assemble the esearch URL
$base = 'https://eutils.ncbi.nlm.nih.gov/entrez/eutils/';
$url = $base . "esearch.fcgi?db=$db&term=$query&usehistory=y";
#post the esearch URL
$output = get($url);
#parse WebEnv and QueryKey
\  \  = 1 if (\output =~/<WebEnv>(\S+)<\backslashWebEnv>/);
$key = $1 if ($output =~ /<QueryKey>(\d+)<\/QueryKey>/);
### include this code for ESearch-ESummary
#assemble the esummary URL
$url = $base . "esummary.fcgi?db=$db&query_key=$key&WebEnv=$web";
#post the esummary URL
$docsums = get($url);
print "$docsums";
### include this code for ESearch-EFetch
#assemble the efetch URL
$url = $base . "efetch.fcgi?db=$db&query_key=$key&WebEnv=$web";
$url .= "&rettype=abstract&retmode=text";
#post the efetch URL
$data = get($url);
nrint "¢data".
```

I used to same way to search for radiofrequency ablation"[MeSH Terms] AND Clinical Trial[ptyp] Here is the url I used:

https://eutils.ncbi.nlm.nih.gov/entrez/eutils/esearch.fcgi?db=pubmed&term=%22radiofrequen cy%20ablation%22[mesh]+AND+%22Clinical%20Trial%22[ptyp]

then the output is XML format, which has exactly the same number of ID as we search in the website.

This XML file does not appear to have any style information associated with it. The document tree is shown below.

```
▼<eSearchResult>
    <Count>2535</Count
    <RetMax>20</RetMax>
    <RetStart>0</RetStart>
  ▼<IdList>
      <IdL18t>
<Id>30392525</Id>
<Id>30382891</Id>
<Id>30320138</Id>

      <Id>30285795</Id>
      <Id>30229917</Id>
      <Id>30229917</Id>
<Id>30143031</Id>
<Id>30111304</Id>
<Id>30075596</Id>
</Id>
      <Id>30040441</Id>
      <Id>29993130</Id>
      <Id>29993130</Id>
<Id>29931084</Id>
<Id>29921526</Id>
<Id>29871374</Id>
      <Id>29847426</Id>
      <Id>29788847</Id>
      <Id>29788847/1d>
<Id>29721503</Id>
<Id>29688123</Id>
<Id>29675772</Id>
      <Id>29663464</Id>
      <Id>29534384</Id>
 "<TermSet>
  <TermSet>
  <TermSet>
  <TermSet>
  <Field>tptyp</Field>
  <Count>818499</Count>
        <Explode>Y</Explode>
      </TermSet>
   </Pre></P>
</Pre>
  ▼<QueryTranslation>
      "radiofrequency ablation"[mesh] AND "Clinical Trial"[ptyp]
   </QueryTranslation>
```

Use the first ID as example, we can use summary to get summary of each ID on the list, using the following url:

https://eutils.ncbi.nlm.nih.gov/entrez/eutils/esummary.fcgi?db=pubmed&id=30392525

This XML file does not appear to have any style information associated with it. The document tree is shown below

```
</
```

Useful tutorial video: Learn more:

Where to learn more?

https://www.ncbi.nlm.nih.gov/books/NBK25501/

YouTube Videos:

2012 (4 min): https://www.youtube.com/watch?v=BCG-M5k-gvEMORE(VIDEOShttps://www.youtube.com/watch?v=iCFVVexp30o

https://ncbiinsights.ncbi.nlm.nih.gov/2017/11/14/five-new-videos-on-youtube-apis-pubmed-refseq-more/