## Working with biological sequence data

# We will use the Biopython package http://biopython.org



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#### Biopython

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#### Introduction

Biopython is a set of freely available tools for biological computation written in Python by an international team of developers.

It is a distributed collaborative effort to develop Python libraries and applications which address the needs of current and future work in bioinformatics. The source code is made available under the Biopython License, which is extremely liberal and compatible with almost every license in the world.

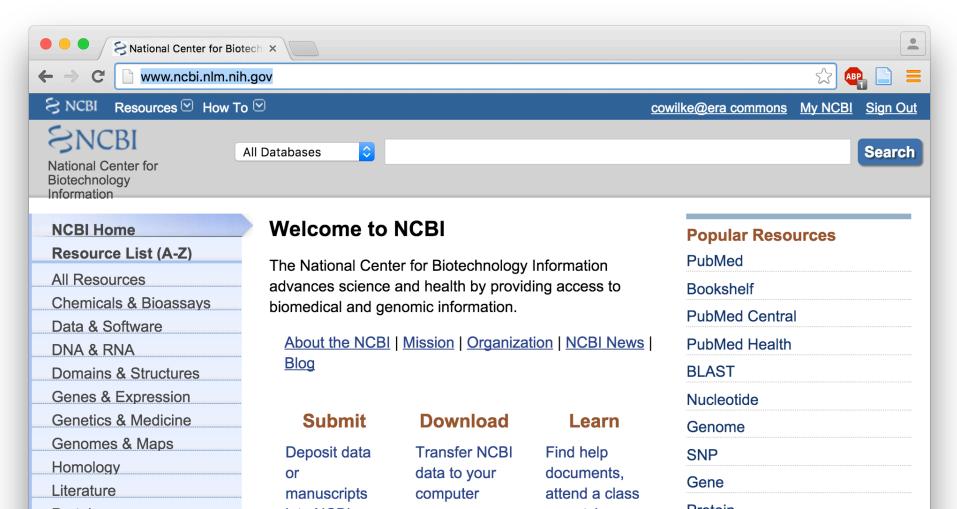
We are a member project of the Open Bioinformatics Foundation (OBF), who take care of our domain name and hosting for our mailing list etc. The OBF used to host our development repository, issue tracker and website but these are now on GitHub.

This wiki will help you download and install Biopython, and start using the libraries and tools.

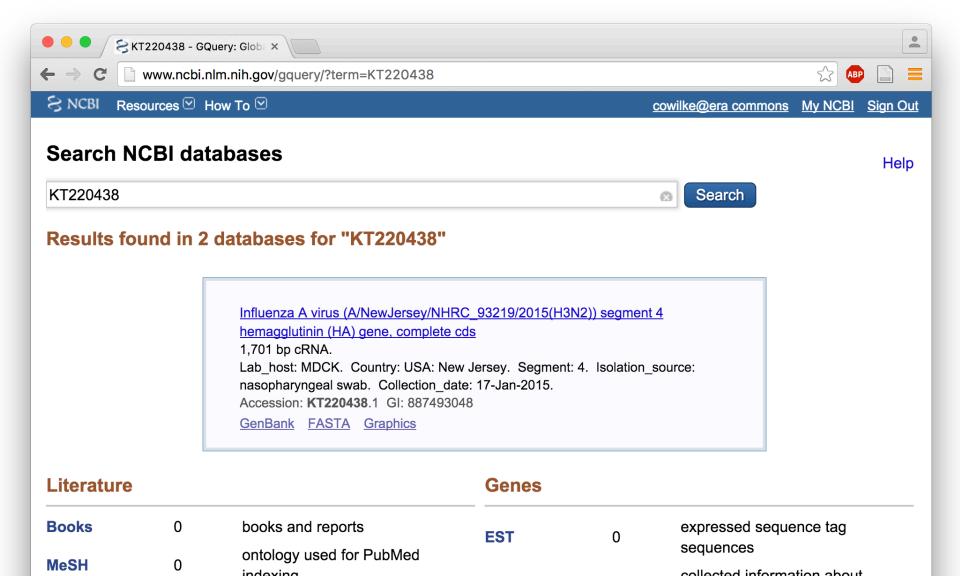
Get Started	Get help	Contribute
Download Biopython	Tutorial (PDF)	What's being worked on
Installation help (PDF)	Documentation on this wiki	Developing on Github
	Cookbook (working examples)	Google Summer of Code

#### Getting biological data: The NCBI databases

http://www.ncbi.nlm.nih.gov/



#### Try search for "KT220438"



#### Direct link to search results

http://www.ncbi.nlm.nih.gov/gquery/?term=KT220438

## A genbank record is just a simple text file

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LOCUS
            KT220438
                                     1701 bp
                                                CRNA
                                                        linear VRL 20-JUL-2015
            Influenza A virus (A/NewJersey/NHRC 93219/2015(H3N2)) segment 4
DEFINITION
            hemagglutinin (HA) gene, complete cds.
ACCESSION
            KT220438
VERSION
            KT220438.1 GI:887493048
KEYWORDS
SOURCE
            Influenza A virus (A/New Jersey/NHRC 93219/2015(H3N2))
            Influenza A virus (A/New Jersey/NHRC 93219/2015(H3N2))
  ORGANISM
            Viruses; ssRNA viruses; ssRNA negative-strand viruses;
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            1 (bases 1 to 1701)
REFERENCE
            Sitz, C.R., Thammavong, H.L., Balansay-Ames, M.S., Hawksworth, A.W.,
  AUTHORS
            Myers, C.A. and Brice, G.T.
  TITLE
            GEISS Influenza Surveillance Response Program
            Unpublished
  JOURNAL
            2 (bases 1 to 1701)
REFERENCE
            Sitz, C.R., Thammavong, H.L., Balansay-Ames, M.S., Hawksworth, A.W.,
  AUTHORS
            Myers, C.A. and Brice, G.T.
            Direct Submission
  TITLE
            Submitted (29-JUN-2015) Operational Infectious Diseases, Naval
  JOURNAL
            Health Research Center, 140 Sylvester Rd., San Diego, CA 92106, USA
            ##Assembly-Data-START##
COMMENT
            Sequencing Technology :: Sanger dideoxy sequencing
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                     Jersey/NHRC 93219/2015(H3N2))"
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                     /strain="A/NewJersey/NHRC 93219/2015"
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/coro+wno-"#3N2"

```
Location/Oualifiers
FEATURES
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                     /codon start=1
                     /product="hemagglutinin"
                     /protein id="AKQ43545.1"
                     /db xref="GI:887493049"
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                     ITNDRIEVTNATELVONSSIGEICDSPHOILDGENCTLIDALLGDPOCDGFONKKWDL
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                     SSFFSRLNWLTHLNYTYPALNVTMPNNEOFDKLYIWGVHHPGTDKDQIFLYAQSSGRI
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                     RSGKSSIMRSDAPIGKCKSECITPNGSIPNDKPFQNVNRITYGACPRYVKHSTLKLAT
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