Working with biological sequence data

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



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Biopython Source Code -

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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.

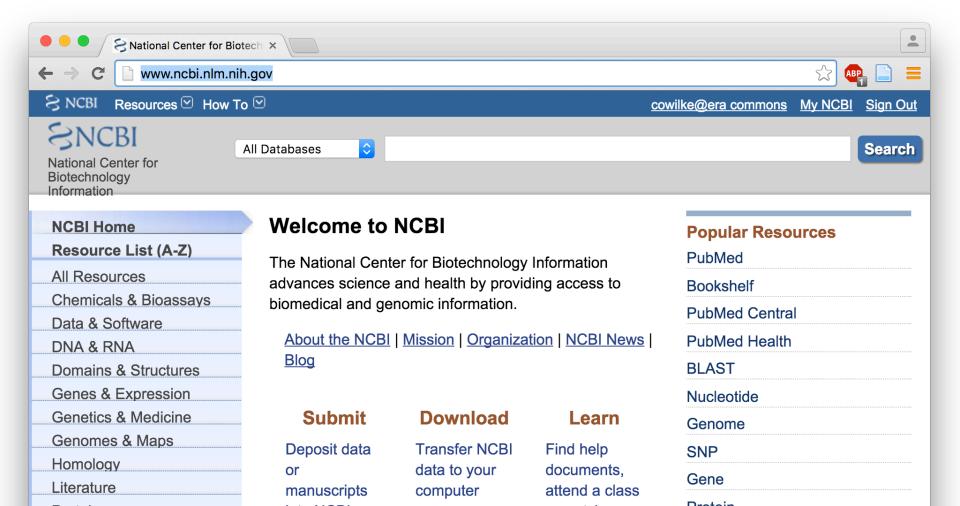
We are a member project of the Open Bioinformatics Foundation (OBF), who take care of our domain name and hosting for our mailing list etc. The OBF used to host our development repository, issue tracker and website but these are now on GitHub.

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

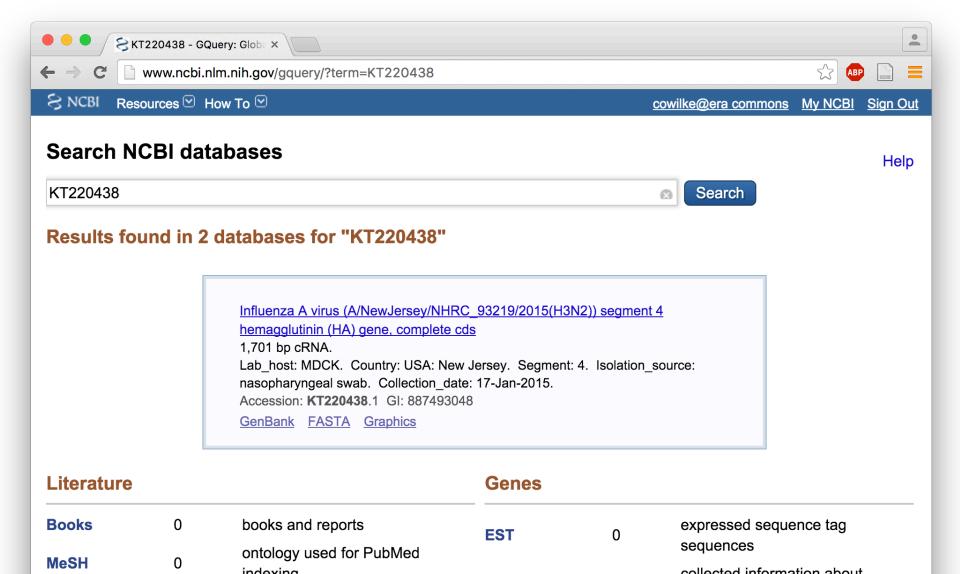
| Get Started | Get help | Contribute |
|-------------------------|-----------------------------|------------------------|
| Download Biopython | Tutorial (PDF) | What's being worked on |
| Installation help (PDF) | Documentation on this wiki | Developing on Github |
| | Cookbook (working examples) |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



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            KT220438
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REFERENCE
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  AUTHORS
            Myers, C.A. and Brice, G.T.
            GEISS Influenza Surveillance Response Program
  TITLE
  JOURNAL
            Unpublished
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            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
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