

# 2020 NCBI CoLab Dataset Survey

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Public reporting burden for this collection of information is estimated to average 2 minutes per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. An agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a current valid OMB control number.

Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to NIH, Project clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0648). Do not return the completed form to this address.

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Page Break

Q1 Select a professional category that best describes you.

- Life Science Researcher
  - Clinical Testing Laboratory Staff
  - Health Professional / Clinician
  - Bioinformatics Professional
  - Computer Scientist / Software Developer
  - Educator
  - Student
  - Librarian / Information Specialist
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Q2 Select a category that best describes your organization.

- College or University
  - Commercial or Industry
  - Non-Profit Organization
  - Government
  - Clinical diagnostics
  - Clinical research
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Q3 What is your preferred way of accessing sequence data? Select one.

- Command line
  - API
  - Web interface
  - Other (please list below) \_\_\_\_\_
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Q4 Which programming languages do you use?

- Python
  - R
  - C++
  - Perl
  - JavaScript
  - Other (please list below) \_\_\_\_\_
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Q5 What type of data are you looking for?

- Gene sequence
  - Gene expression
  - Genome sequence
  - Genetic variation
  - Associated metadata for sequences
  - Other (please list below)
- 

Q6 How do you query for data?

- Taxonomy
  - Gene symbol
  - NCBI accession
  - Ensembl accession
  - Other (please list below)
-

Q7 What metadata format(s) do you prefer for your workflow? Select all that apply.

- CSV
  - TSV
  - JSON
  - YAML
  - Other \_\_\_\_\_
- 

Q8 What annotation format(s) do you prefer for your workflow? Select all that apply.

- GFF
  - GTF
  - Other (please list below)  
\_\_\_\_\_
- 

Q9 Do you have any recommendations on how we can improve NCBI Datasets?

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Q10 On a scale from 0-10, how likely are you to recommend NCBI Datasets to a friend or colleague?

- 0
- 1
- 2
- 3
- 4
- 5
- 6
- 7
- 8
- 9
- 10

End of Block: Default Question Block

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