

NCBI Gene User Survey

We want to improve NCBI's products and services. We would like to learn more about how this site helps you with your work, and what we can do better. Please click "next" below to get started.

OMB Control Number: 0925-0648

Expiration Date: 05/31/2021

Public reporting burden for this collection of information is estimated to average 5 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.

[Next](#)

[Privacy & Cookie Policy](#)

1. Which professional category best describes you? Please select only one.

- Bioinformatics professional
 Educator
 Healthcare professional
 Librarian / Information Specialist
 Life Science Researcher
 Student
 Technician
 Computer Scientist / Software Developer
 Other (please specify)

2. Please pick one category that best describes your organization.

- College or University
 Commercial / Industry
 Non-profit Organization
 Government
 Other (please specify)

3. How often do you use NCBI Gene?

- Every day
 A few times a week
 About once a week
 A few times a month
 Once a month
 Less than once a month

4. How do you obtain gene-related information from NCBI? Please check all that apply.

- Gene web pages
 Gene FTP site
 NCBI programming utilities APIs (E-Utilities, EDirect)
 Other (please specify)

5. Which of the following organism(s) are the focus of your research? Please check all that apply.

- Human
 Mouse
 Mammals other than mouse or human
 Non-mammalian vertebrates
 Invertebrates
 Plants
 Fungi
 Protozoa
 Prokaryotes
 Viruses

6. What was the purpose of your visit to Gene today? Please check all that apply.

- Find general information about a gene
 Find sequences
 Use links to other resources (GEO, SNP, ClinVar, etc.)
 Get information about gene expression
 Get information about gene / protein function
 Find publications
 Other (please specify)

7. What sections or features of the NCBI Gene page do you use in your work? Please check all that apply.

- Nomenclature
 Summary
 Graphical View/Sequence Viewer
 Link to Gene sequence and location
 Expression
 Interactions and Pathways
 Orthologs
 Variation
 Genomic context
 Publications
 RefSeq transcripts and proteins
 Gene Ontology

8. What publication information would you like to access from NCBI Gene? Please check all that apply.

- All publications
 The most recent publications
 Selected publications that provide information about gene function
 GeneRIF summaries of functional information about the gene
 I don't use any of the publication information in Gene

9. How do you use the information you obtain from NCBI Gene? Please check all that apply.

- Experiment/assay design
 Manuscript preparation
 Clinical analysis
 Teaching
 Generating reports/protocols
 Other (please specify)

10. What content do you use from the NCBI Gene FTP files? Please check all that apply.

- RefSeq-Ensembl matches from gene2ensembl.gz and gene2vega.gz
 Gene Ontology from gene2go.gz
 I don't use the Gene FTP site
 Gene summary information from gene_info.gz
 Genome annotation from ASN_BINARY
 RefSeq accessions for each gene from gene2refseq.gz
 RefSeq-UniProtKB matches from gene_refseq_uniprotkb_collab.gz
 Publications from gene2pubmed.gz
 Summaries from ASN_BINARY
 Orthology data from gene_orthologs.gz
 GeneRIF
 Other

11. Which of the following additions to NCBI Gene would benefit your research the most? Please check all that apply.

- Expanded orthology information to include organisms outside the vertebrates
 Expression data for more organisms
 Expression data from more tissues/developmental stages
 Comparison between RefSeq annotation and Ensembl or GenBank
 Expanded reporting of UniProt accessions (both SwissProt and TrEMBL)
 Advanced search for expression data
 Pan-genome analyses
 Transcript variant-specific expression information
 Sequence data from multiple strains/breeds/cultivars
 Other (please specify)

12. Did you find what you were looking for today?

- Yes
 No

13. How easy was it to find what you were looking for on the NCBI Gene website?

- Extremely easy
 Very easy
 Somewhat easy
 Not so easy
 Not at all easy

14. Please provide any additional suggestions you may have on improving NCBI Gene.

15. How likely is it that you would recommend NCBI Gene to a friend or colleague?

Not at all likely

Extremely likely

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

Please write to us at info@ncbi.nlm.nih.gov you'd like to provide additional feedback about NCBI Gene resources.

Prev

Done