

NCBI Protein Resources 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.

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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.

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

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

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

- College or University
- Commercial / Industry
- Non-profit Organization
- Other (please specify)
- Research Institute
- Government
- Hospital/health care organization

3. What are your areas of research? Please check all that apply.

- Biochemistry & Molecular Biology
- Bioinformatics/Computational Biology
- Genetics
- Microbiology
- Immunology
- Medicine
- Chemistry & Pharmacology
- Other (please specify)

4. What task are you trying to accomplish at the NCBI site today?

5. What kind of information do you want to find? Please check all that apply.

- Sequence(s) for genes, transcripts, proteins, isoforms, alignments
- Nomenclature, official name/symbol, aliases)
- Function
- Physical properties such as 3D/secondary structure, MW, pI, kinetics (Km, v_{max}, Kd), location of motifs/domains/key residues
- Molecular interactions such as protein-protein, protein-ligand, protein-nucleotide, pathways
- Experimental study data such as activity assays, expression levels/patterns, proteomics.
- Reagents, for example, purified protein, ligands/substrates/modulators, antibodies, expression clones, drugs, inhibitors, molecular probes
- Genomic context: location in the genome and the genes/regulatory regions are located near your gene of interest
- Literature
- Clinically-related information including disease/disorder/condition or phenotype, genetic variations
- Other (please specify)

6. Please outline your workflow that would include the data you are seeking today at NCBI. Please tell us about other NCBI resources as well as non-NCBI resources that you use in your processing.

7. Please tell us about any information you need but can not find or tasks that you can not perform at NCBI.

8. Which of the following issues limit your effective use of NCBI's current protein resources? Please choose all that apply.

- I don't know how to/where to look for the information I need.
- I can't find what I am looking for (no relevant results).
- I am overwhelmed by the number of results. The relevant results are buried.
- The results are misleading.
- Search and retrieval services are too slow.
- Search and retrieval services don't always work from my location.
- Other (please specify)

9. How likely is it that you would recommend NCBI protein resources to a friend or colleague?

Not at all likely

Extremely likely

0	1	2	3	4	5	6	7	8	9	10
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We welcome your input! Please contact us at suggest@ncbi.nlm.nih.gov if you would like to provide additional feedback about NCBI protein resources.

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Done