

NCBI Sequence Search 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 3 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

1. Which professional categories best describe you? Please check all that apply.

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

2. Which of the following categories best describes your organization?

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

3. What were you looking for today at NCBI? Please check all that apply.

- Gene report
- Gene sequence
- Representative transcript sequence
- Set of transcript sequences
- Representative protein sequence
- Set of protein sequences
- Genome assembly
- Set of assemblies
- Other (please specify)

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

- Yes
- No

5. How easy was it to find what you were looking for on our website?

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

6. What suggestions do you have for improving this search service?

7. How likely is it that you would recommend this service 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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8. May we contact you to get additional feedback?

- Yes
- No

9. Please enter your email address below.

Prev

Done