

NCBI BLAST 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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NCBI BLAST Survey

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. How frequently do you use each of the following services?

	Daily	Weekly	Monthly	Less often than monthly	Never
BLAST webpages other than the search forms (help documents etc.)	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Nucleotide BLAST (blastn)	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Protein BLAST (blastp)	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Translated BLAST (blastx, tblastn, tblastx)	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
BLAST Help, News, and documentation webpages	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

10. Please indicate how strongly you agree or disagree with the following statements about the NCBI BLAST web service.

	Strongly disagree	Disagree	Neither Agree nor Disagree	Agree	Strongly agree	N/A
BLAST services are easy to find from the BLAST homepage.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Knowing which BLAST databases to choose is easy.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Finding BLAST databases for a specific organism is easy.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The default settings on the BLAST search pages work well for what I do.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
BLAST results are returned quickly.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
BLAST results are easy to read and understand.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The Graphic Summary of the results is useful.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The Descriptions table has a clear and understandable layout.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The results download capabilities are easy to use and provide what I need.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The Graphics (Graphical Sequence Viewer) links from the results are a valuable resource.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
The other reports (Taxonomy, Distance Tree of Results, etc.) are useful.	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

11. Please provide any specific examples of any negative experiences you have had with NCBI BLAST products.

12. How likely is it that you would recommend NCBI BLAST 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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13. If there was one thing you could change about BLAST, what would it be?

14. Would you be willing to pay for the ability to run large numbers of BLAST searches with fewer restrictions on use and higher throughput?

Yes

No

15. Where do you get your information or news about BLAST?

BLAST Web pages / Help documents

BLAST announce emails / newsletter

Conference presentations

Journals- printed/online

NCBI Insights Blog or other Blogs

Social media (Twitter, FaceBook, LinkedIn)

NCBI or other YouTube videos

Discussion groups/forums (SeqAnswers, BioStars etc.)

Work colleagues

Other (please specify)

16. Which of these other sequence analysis products do you use?

BLAT

BWA

SOAP

Bowtie

HMMER

PLAST

ExPasy - SIB BLAST

BlastStation

EBI-EMBL Tools

Geneious

Lasergene – DNASTAR

CLC Genomics Workbench – QIAGEN

MacVector

UGENE

Galaxy

None

Other (please specify)

17. Do you have a need to run NCBI's BLAST on your own server(s) or on those provided by a third party?

Yes

No

If you use a 3rd party provider, which one do you use? For example: Amazon Web Services (AWS), Google Cloud Platform Engine, Microsoft Azure, IBM Cloud, Rackspace, other – please specify

18. May we contact you to get additional feedback?

Yes

No

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