

Note: The attached survey splits at question 3 on page 2. Those who have submitted data before and select any of the first nine choices on question 3 go to the questions (4-11) on page 3. Those who have not successfully submitted and select any of the last three choices answer the questions (4-10) on page 4.

NCBI Submission 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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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. Have you ever submitted sequence data to NCBI via the products below? Please check all that apply or indicate that you have not submitted.

- BankIt
- SRA
- BioProject
- BioSample
- TSA
- WGS
- GEO
- Sequin
- Submitted but don't remember product
- Started but did not complete submission
- Have never tried to submit sequences
- Other (please specify)

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4. How easy was it to submit your sequence data? If you have made multiple submissions, please evaluate your most recent submission.

- Very easy
- Easy
- Neither easy nor difficult
- Difficult
- Very difficult

5. Why did you select the answer you gave in the previous question?

6. How did you find out what to do the first time you submitted sequence data? Please check all that apply.

- Google or other search engine
- NCBI (www.ncbi.nlm.nih.gov) or NIH home page (www.nih.gov) search
- Submission portal page (https://submit.ncbi.nlm.nih.gov/)
- Followed link from an academic, industry or government website
- Other (please specify)

7. What did you like least/most about your sequence submission experience?

8. How easy was it for you to complete the following sequence submission-related tasks?

	Very easy	Easy	Neither easy nor difficult	Difficult	Very Difficult	Didn't do
Identify correct submission tool	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Complete NCBI login	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Begin new submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Complete a previous submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Understand and gather required materials for submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Follow instructions and enter data on forms	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Enter taxonomic or other data specific to your submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Upload files	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Correct form errors during submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Use NCBI help materials to complete submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Understand submission status, steps, and timeline	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

9. How can we improve sequence submissions for you?

10. What was your primary goal when you visited the site today?

- Prepare in advance to make a new data submission
- Begin new data submission & submit to NCBI
- Complete existing data submission & submit to NCBI
- Request a change to previously-submitted data
- Request an update on the status of my submission or accession number
- Assist a friend or colleague with their submission
- Other (please specify)

11. How likely is it that you would recommend the NCBI Submission Portal 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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4. To what NCBI sequence submission product(s) do you plan to submit? Please check all that apply or indicate that you are unsure or will not submit.

- BankIt
- SRA
- BioProject
- BioSample
- TSA
- WGS
- GEO
- Sequin
- Unsure
- Will not submit
- Other (please specify)

5. What was your primary goal when you visited the site today?

- Prepare to make a data submission later
- Begin new data submission & submit to NCBI
- Complete existing data submission & submit to NCBI
- Assist a friend or colleague with their submission
- Other (please specify)

6. What, if anything, is currently preventing you from submitting sequence data to NCBI? Please check all that apply.

- Scientific research not yet complete
- Not yet sure how to submit this data
- Awaiting data or input from a colleague
- Errors in my submission so far
- Need NCBI guidance
- Nothing
- Other (please specify)

7. Which of the following website features do you think would help you begin your submission to NCBI? Please rank any features you think would be useful.

	Very Useful	Useful	Neutral	Not Useful	Unsure	No opinion
Diagram illustrating step-by-step submission process	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Examples of successful submissions	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Option to select sequence data type, task (submit, edit) and be guided through submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Required materials checklist	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Option to upload sequence data and see your submission automatically routed to correct place	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Ability to search FAQs on submission steps	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Centralized list of file templates for submission, by product	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Submission tips	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

8. How can we improve sequence submissions for you?

9. Which of the following words do you think accurately describe NCBI's sequence submission information online? Please check all that apply.

- Not enough detail
- Too much detail
- Easy to understand
- Difficult to understand
- Helped me begin my submission
- Did not help me begin my submission
- Too much text
- Too many diagrams and charts
- Clear terminology
- Confusing terminology
- None of these apply

10. How likely is it that you would recommend the NCBI Submission Portal 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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The NCBI team values our users and welcomes all input! Please write to us at suggest@ncbi.nlm.nih.gov if you'd like to share additional feedback or help us test new sequence submission features.

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Done