

OMB Control Number: 0925-0648 Expiration Date: 05/31/2021

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

- Bioinformatics Professional
 Life Science Researcher
 Computer Scientist / Software Developer
 Student
 Educator
 Other (please specify)

2. Pick one category that best describes your organization.

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

3. Are you submitting a single sequence or a set of sequences?

- Single sequence
 Set of sequences

4. How many times have you submitted to GenBank using BankIt?

- 1-4
 5-50
 50+

5. How easy was it to complete your submission? If you have made multiple submissions, please evaluate your most recent one.

- Very easy
 Easy
 Neither easy nor difficult
 Difficult
 Very difficult

6. Explain why you chose the answer provided in the previous question.

7. What type of data do you typically submit?

- mRNA
 Eukaryotic genomic or organellar DNA with coding regions
 Prokaryotic genomic DNA
 Virus sequences
 Complete organelle genomes
 Phylogenetic sets of marker genes
 Other (please specify)

8. How do you organize your sequence data?

- Gene, locus, or region
 Organism type
 Genomic location (organelle, plasmid)
 Other (please specify)

9. How easy was it to complete the following submission tasks?

	Very easy	Somewhat easy	Neither easy nor difficult	Somewhat difficult	Very difficult	N/A
Understanding and gathering required materials for submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Following instructions and entering data on forms	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Entering 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"/>
Uploading files	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Annotating features (Adding CDS, gene, etc.)	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Downloading the flat file	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Propagating features	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Correcting web form errors during submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Using NCBI help materials to complete submission	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Understanding submission status, steps & timeline	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

10. If you found feature annotation difficult, what were some of the reasons for that?

- Complexity of the web application
 Complexity of the biology involved
 Complexity of the web application and biological complexity
 I had no difficulties with annotation
 Other (please specify)

11. Please suggest how we can make completing feature annotation easier for you.

12. Which of the following improvements would be helpful? Please check all that apply.

- Clearer instructions on web forms
 Improved error messages
 Links to examples
 More diagrams or visual aids
 Improved workflow during submission
 Ability to check the file **format** in advance
 Ability to check **data** in advance
 Other (please specify)

13. What features should NCBI add in the future to help with submission? Please check all that apply.

- Ability to update my submission in the same place I submit data
 Additional file formats accepted (ASN1, other)
 Ability to submit as part of a group
 Ability to manage submission group(s)
 Ability to share submission data with collaborators or publisher/reviewer
 Other (please specify)

14. Please suggest ways to make submission to GenBank easier.

15. How likely is it that you would recommend this site 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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Please write to us at suggest@ncbi.nlm.nih.gov if you would like to provide additional feedback on this site.

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Done