

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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Done