



The “Organism” Page

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This page will appear after the “Nucleotide” page only if you made an error in the organism FASTA format or did not provide an organism in the FASTA definition line at all. If you correctly entered an organism in the definition line for each sequence you are submitting in the “Nucleotide” page, the “Organism” page will not appear, and you will go directly to the next page of the form.

If you have a single sequence, or multiple sequences from the same organism:

1. Begin entering the organism name in the text box to the right of “Organism name”:
2. Once you begin entering the organism name in the text box provided, BankIt will provide a list of species names based on the text you enter (Figure 10).
3. Select the appropriate organism name for your submission. If you do not see the appropriate organism listed, type the correct and complete organism name into the text box.
4. Click the “Continue” button.
5. You will see the organism name appear with its sequence ID in a static table below “Continue” button. Check to be sure the organism you selected or typed in is correct.
6. Click on the “Continue” button again to go to the next page of the submission form.

If you have multiple sequences with multiple names:

1. Select an organism file from your local computer using the “Browse” button to the right of the words “Upload File”. See the BankIt Help documentation for [instructions](#) for creating an organism file.
2. Click the “Continue” button.
3. You will see the organism names and sequence IDs from your organism table appear in a static table below “Continue” button. Check to be sure the organism names and sequence IDs are correct.
4. Click on the “Continue” button again to go to the next page of the submission form.

GenBank Submissions

Contact Reference Nucleotide **Organism** Set/Batch Submission Category Source Modifiers Primers Features
Review and Correct

Submission # 1449155

Fill in missing Organism information

You did not include the name of the organism from which the sequence was isolated. Please enter the organism name below. (For future sequence submissions, be sure to use the [FASTA](#) format.)

Organism Name Input same organism name for all sequences.

Upload File In [Organism](#) file?

Continue

Organism

Abbott's mouse (*Mus abbotti*)
 Aceramarca gracile mouse opossum (*Gracilinanus aceramarcae*)
 Afghan mouse-like hamster (*Calomyscus fulvipes*)
 Agile Gracile Mouse Opossum (*Gracilinanus agilis*)
 Allen's wood mouse (*Hylomyscus alleni*)
 Alston's brown mouse (*Scotinomys teguina*)
 Andean big-eared mouse (*Auliscomys subandinus*)
 Andean pygmy mouse (*Eligmodontia puerulensis*)

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Begin entering the organism name for your sequence, and BankIt will provide a list of species names based on the text you enter.

Review the species names provided to find the organism that the sequence came from. Click on it to enter it into the text box. If the organism is not included in this list, continue typing the complete and correct organism name.

Figure 10: The Organism page showing a species drop down list released as you enter text in “Organism Name” text box. Figure text gives information about the drop-down list of species names.