

Filter out Short Sequences

To use this workflow input a set of sequences and set a minimum sequence length. All sequences with length less than the specified value will be filtered out. The result will be written into a FASTA file by default.



How to Use This Sample

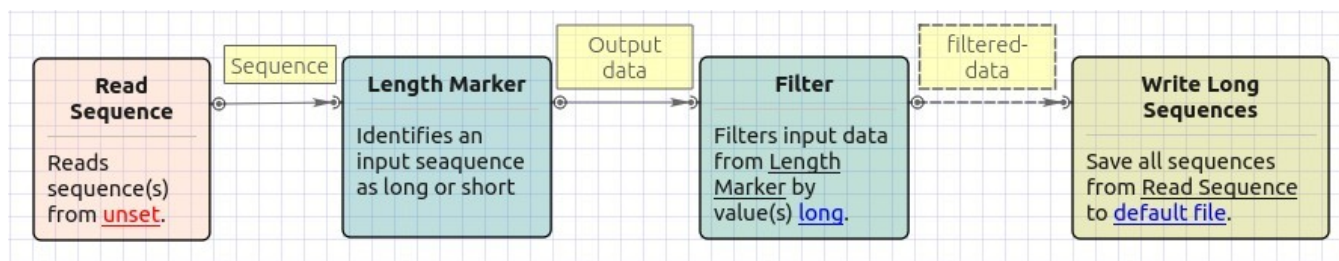
If you haven't used the workflow samples in UGENE before, look at the "[How to Use Sample Workflows](#)" section of the documentation.

Workflow Sample Location

The workflow sample "Filter out Short Sequences" can be found in the "Scenarios" section of the Workflow Designer samples.

Workflow Image

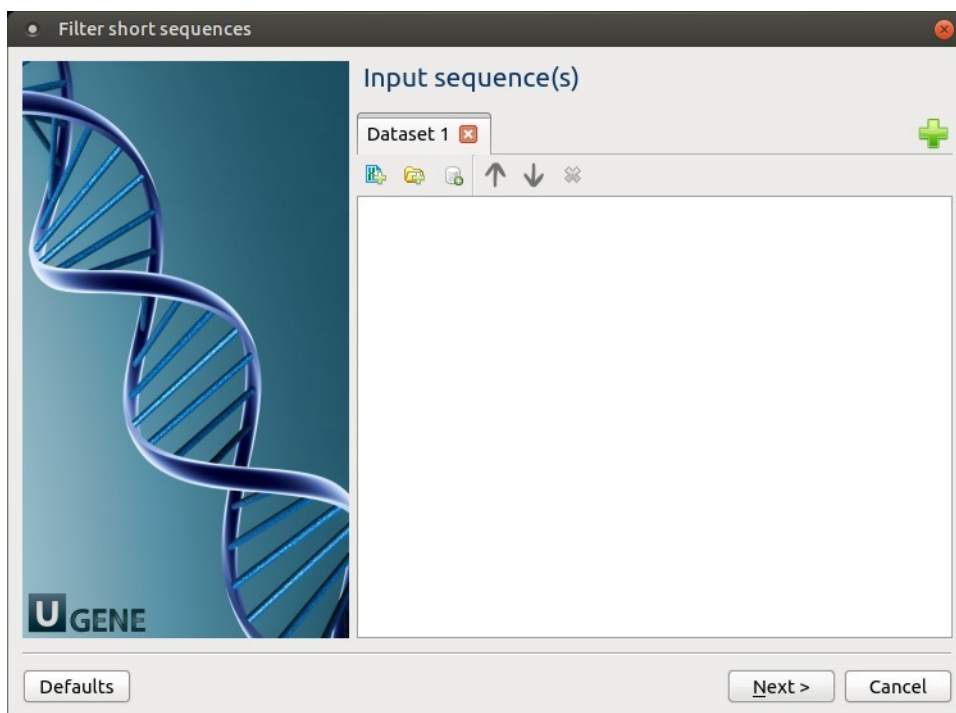
The opened workflow looks as follows:



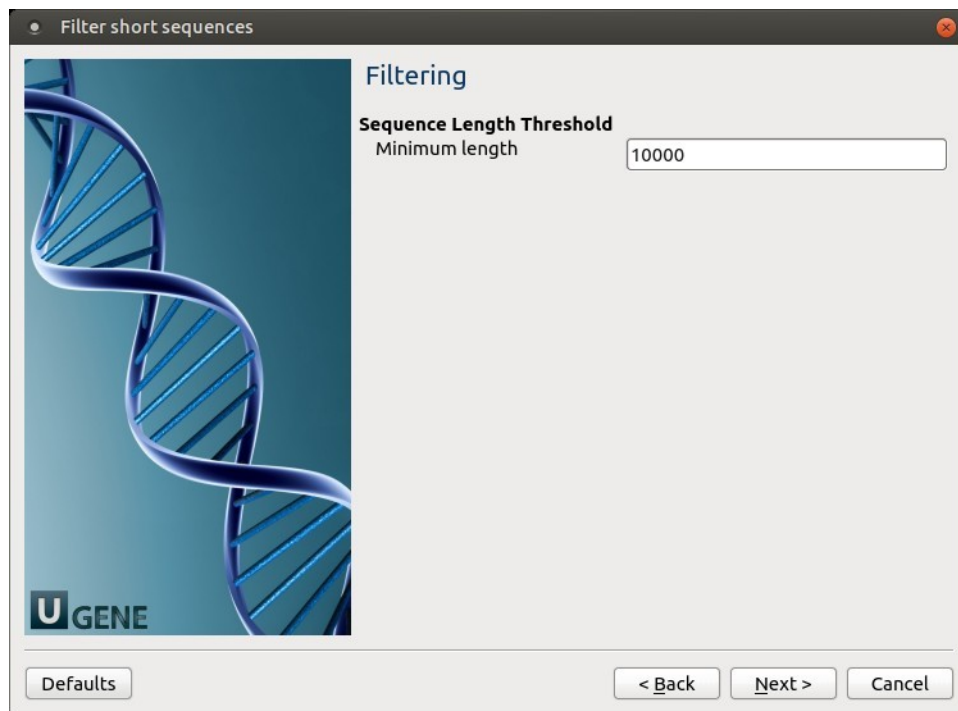
Workflow Wizard

The wizard has 3 pages.

1. Input sequence(s): On this page, input files must be set.



2. Filtering: The filtering parameters can be changed here.



The following parameters are available:

Minimum length	Minimum sequence length
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3. Output data: On this page, the output file can be selected:

