

# 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

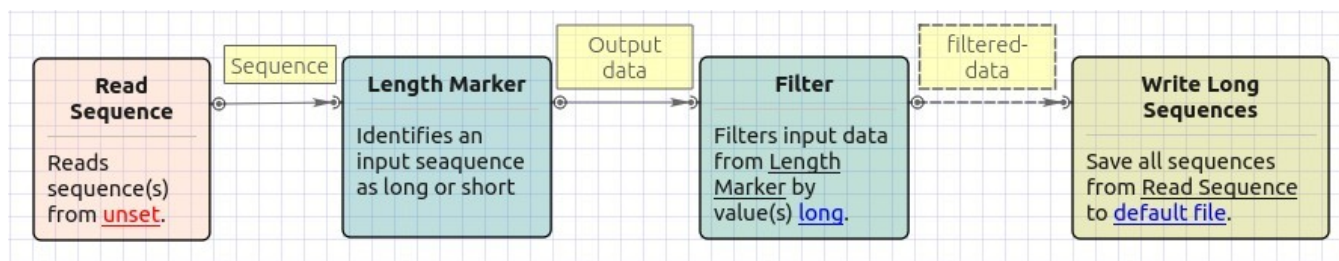
If you haven't used the workflow samples in UGENE before, look at the "[How to Use Sample Workflows](#)" section of the documentation.

## Workflow Sample Location

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

## Workflow Image

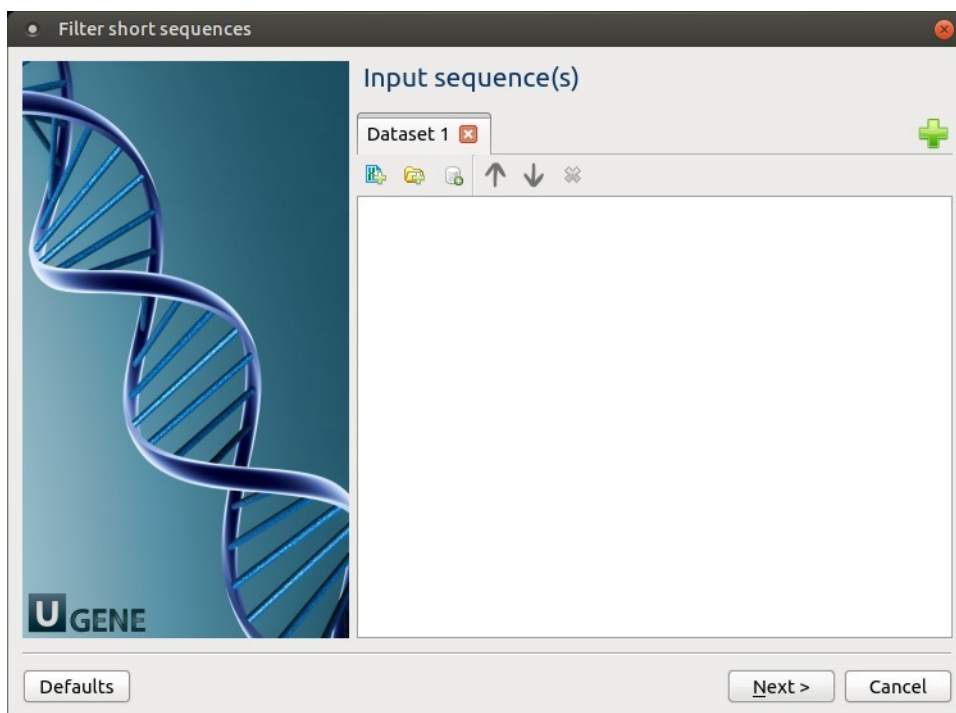
The opened workflow looks as follows:



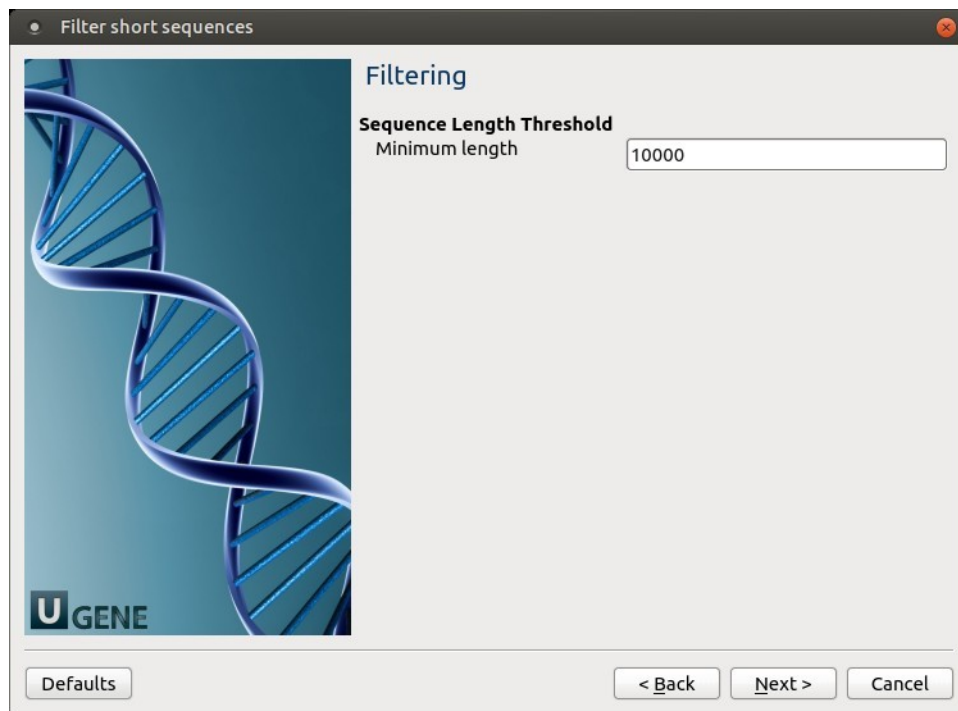
## Workflow Wizard

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:

