Get Amino Translations of a Sequence

The workflow takes a nucleotide sequence as input and returns two files: translations of the sequence and translations of the complement sequence.

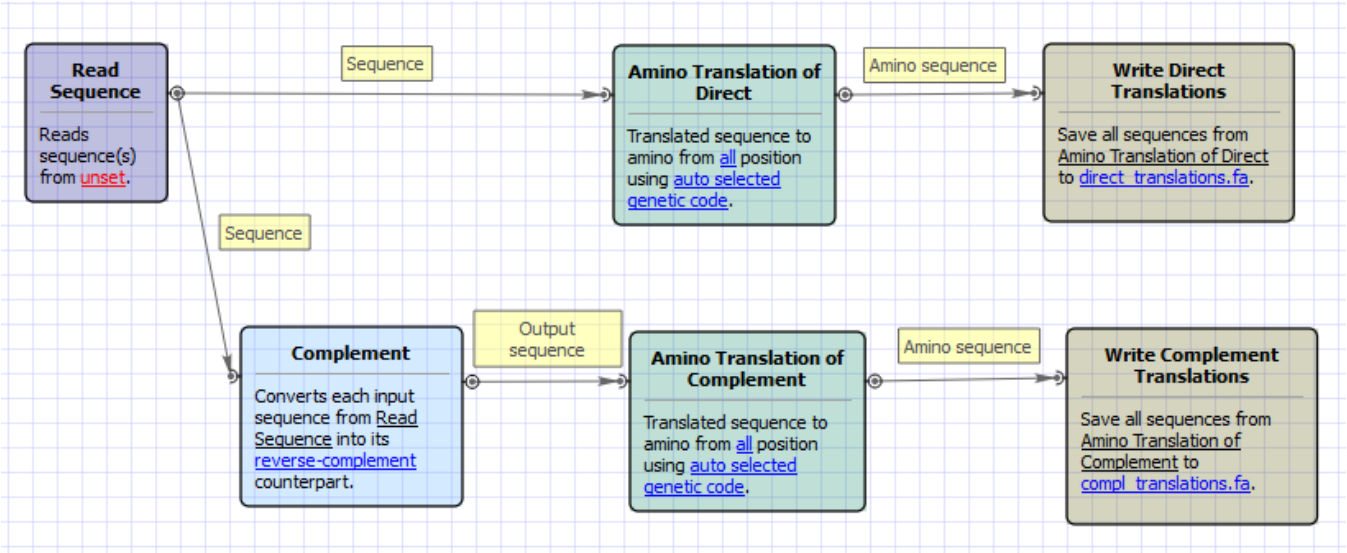
> **How to Use This Sample**
>
> If you haven't used the workflow samples in UGENE before, look at the "How to Use Sample Workflows" section of the documentation.

Workflow Sample Location

The workflow sample "Get Amino Translations of a Sequence" can be found in the "Scwnarios" section of the Workflow Designer samples.

Workflow Image

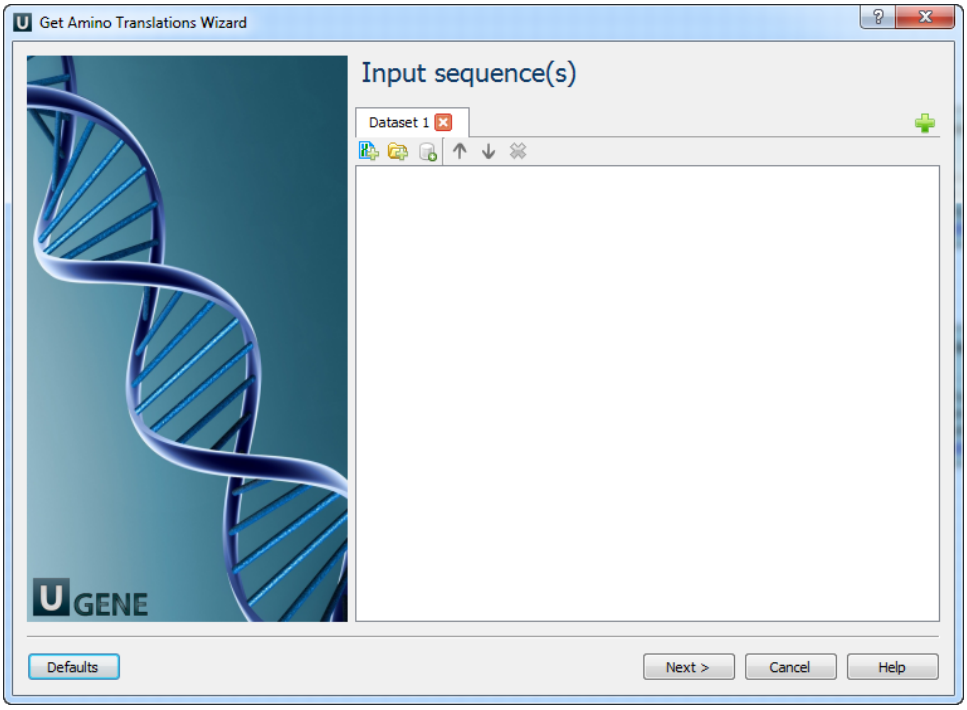
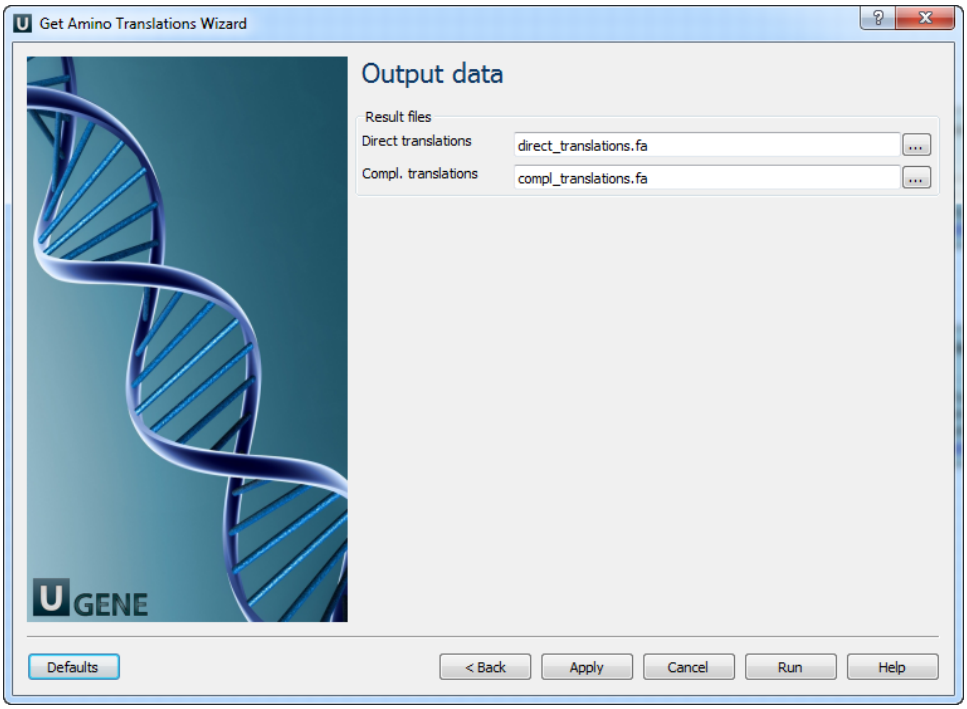
The opened workflow looks as follows:



Workflow Wizard

The wizard has 2 pages.

1. Input sequence(s): On this page you must input sequence(s).



2. Output Data: On this page you can modify result files settings.