

Quality Control by FastQC

FastQC aims to provide a simple way to do some quality control checks on raw sequence data coming from high throughput sequencing pipelines. It provides a molecular set of analyses which you can use to give a quick impression of whether your data has any problems of which you should be aware before doing any further analysis.



How to Use This Sample

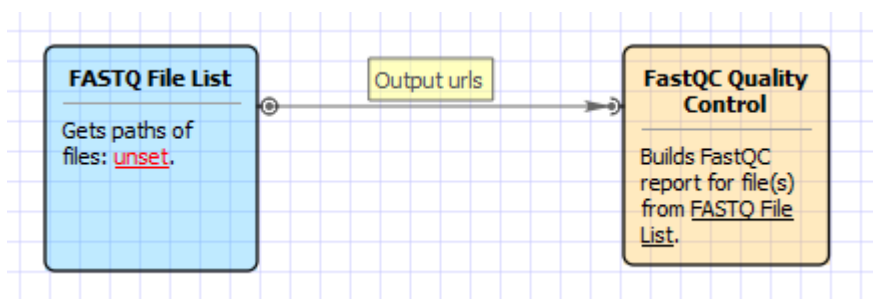
If you haven't used the workflow samples in UGENE before, look at the ["How to Use Sample Workflows"](#) section of the documentation.

Workflow Sample Location

The workflow sample "Quality Control by FastQC" can be found in the "NGS" section of the Workflow Designer samples.

Workflow Image

The workflow is the following:



Workflow Wizard

The wizard has 1 page.

1. High Throughput Sequence QC Report by FastQC: On this page you must input FASTQ file(s) and optionally modify advanced parameters.

