

Merge Annotations Element

Writes all supplied sequences to file(s) in FASTQ format.

Element type: write-fastq

Parameters

Parameter	Description	Default value	Parameter in Workflow File	Type
Output file (required)	Location of the output data file. If this attribute is set, then the "Location" slot is not taken into account.		url-out	<i>string</i>
Existing file	If a target file already exists, you can specify how it should be handled: either overwritten, renamed or appended (if supported by file format).	Rename	write-mode	<i>numeric</i> Available values are: <ul style="list-style-type: none">• 0 - for overwrite• 1 - for append• 2 - for rename
Accumulate objects	Accumulates all incoming data in one file or creates separate files for each input. In the latter case, an incremental numerical suffix is added to a file name.	True	accumulate	<i>boolean</i>

Input/Output Ports

The element has 1 *input port*.

Name in GUI: *Sequence*

Name in Workflow File: in-sequence

Slots:

Slot In GUI	Slot in Workflow File	Type
Sequence	sequence	<i>sequence</i>
Location	url	<i>string</i>