

# **Write Alignment Element**

Writes all supplied alignments to file(s) in selected format.

## Parameters in GUI

Parameter	Description	Default value
<b>Output file</b> (required)	Location of the output data file. If this parameter is set, then the "Location" slot is not taken into account.	
<b>Existing file</b>	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
<b>Document format</b>	Format of the output file.	clustal

## Parameters in Workflow File

Type: write-msa

Parameter	Parameter in the GUI	Type
<b>url-out</b>	<b>Output file</b>	<i>string</i>
<b>write-mode</b>	<b>Existing file</b>	<i>numeric</i> Available values are: <ul style="list-style-type: none"> <li>• 0 - for overwrite</li> <li>• 1 - for append</li> <li>• 2 - for rename</li> </ul>
<b>document-format</b>	<b>Document format</b>	<i>string</i> Available values are: <ul style="list-style-type: none"> <li>• clustal</li> <li>• mega</li> <li>• msf</li> <li>• sam</li> <li>• srfasta</li> <li>• stockholm</li> </ul>

## Input/Output Ports

The element has 1 *input port*:

**Name in GUI:** Multiple sequence alignment

**Name in Workflow File:** in-msa

**Slots:**

Slot In GUI	Slot in Workflow File	Type
<b>MSA</b>	<b>msa</b>	<i>msa</i>
<b>Location</b>	<b>url</b>	<i>string</i>