

Read Alignment Element

Reads multiple sequence alignments (MSAs) from local or remote files.

Parameters in GUI

Parameter	Description	Default value
Input files (required)	Semicolon-separated list of paths to the input files.	

Parameters in Workflow File

Type: read-msa

Parameter	Parameter in the GUI	Type
url-in	Input files	<i>string</i>

Input/Output Ports

The element has 1 *output port*:

Name in GUI: *Multiple sequence alignment*

Name in Workflow File: out-msa

Slots:

Slot In GUI	Slot in Workflow File	Type
MSA	msa	<i>msa</i>
Source URL	url	<i>string</i>