Extract Consensus from Alignment as Sequence

Extract the consensus sequence from the incoming multiple sequence alignment.

Parameters in GUI

Parameter	Description	Default value
Algorithm	The algorithm of consensus extracting.	
Threshold	The threshold of the algorithm.	100
Keep gaps	Set this parameter if the result consensus must keep the gaps.	True

Parameters in Workflow File Type: extract-msa-consensus-sequence

Parameter	Parameter in the GUI	Туре
algorithm	Algorithm	string
threshold	Threshold	numeric
keep-gaps	Keep gaps	boolean

Input/Output Ports The element has 1 input port.

Name in GUI: in-msa

Name in Workflow File: in-msa

Slots:

Slot In GUI	Slot in Workflow File	Туре
MSA	msa	msa

And 1 output port:

Name in GUI: out-sequence

Name in Workflow File: out-sequence

Slots:

Slot In GUI	Slot in Workflow File	Туре
Sequence	sequence	seq