Extract Transcript Sequences with gffread Element

Extract transcript sequences from the genomic sequence(s) with gffread.

Parameters in GUI

Parameter	Description	Default value
Output sequences	The url to the output file with the extracted sequences.	

Parameters in Workflow File

Type: gffread

Parameter	Parameter in the GUI	Type
url-out	**Output sequences**	*string*

Input/Output Ports

The element has 1 *input port*:

Name in GUI: Input transcripts

Name in Workflow File: in-data

Slots:

Slot In GUI	Slot in Workflow File	Type
Genomic sequence url	**genome**	*string*
Transcripts url	**transcripts**	*string*

And 1 *output port*:

Name in GUI: Extracted sequences url

Name in Workflow File: extracted-data

Slots:

Slot In GUI	Slot in Workflow File	Type
sequences	**sequences**	*string*