

Extract Transcript Sequences with gffread Element

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

Element type: gffread

Parameters

Parameter	Description	Default value	Parameter in Workflow File	Type
Output sequences	The url to the output file with the extracted sequences.		url-out	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