

Change Chromosome Notation for VCF Element

Changes chromosome notation for each variant from the input, VCF or other variation files.

Element type: rename-chromosome-in-variation

Parameters

Parameter	Description	Default value	Parameter in Workflow File	Type
Replace prefixes	Input the list of chromosome prefixes that you would like to replace. For example "NC_000". Separate different prefixes by semicolons.		prefixes-to-replace	string
Replace by	Input the prefix that should be set instead, for example "chr".		prefix-replace-with	string

Input/Output Ports

The element has 1 *input ports*:

Name in GUI: Input file URL

Name in Workflow File: in-file

Slots:

Slot In GUI	Slot in Workflow File	Type
Source URL	url	string

And 1 *output port*:

Name in GUI: Output file URL

Name in Workflow File: out-file

Slots:

Slot In GUI	Slot in Workflow File	Type
Produced URL	url	string