



SelectFeaturesRows Documentation

Description: Generate a .res, .gct, .snp, .cn, or .odf file containing only specified features (i.e. rows).

Author: Joshua Gould (Broad Institute), gp-help@broad.mit.edu

Summary: This module generates a file containing only a subset of features represented in the input file. The user specifies the features (i.e. 'rows') to include in the new file, and the order in which they appear, by providing a text file that contains one unique feature identifier per line. Specific columns can be extracted using the *SelectFeaturesColumns* module.

Parameters:

Name	Description
input.filename	input file - .res, .gct, .snp (non-allele-specific), .cn, or odf, file from which the slice is extracted
list filename	text file containing one identifier per line that specifies the rows to select.
output	output filename - .res, .gct, .snp, .cn, .odf

Output Files:

1. output file containing only the specified features (rows)

Platform dependencies:

Module type: GeneListSelection
CPU type: any
OS: any
Language: Java