

gctRowKeep, v13

Filters a gct file to keep a user-defined list of reagents, also filters out NaNs (optional)

Author: Barbara Weir, Broad Institute

Contact: bweir@broadinstitute.org

Algorithm Version:

Introduction

gctRowKeep is a GenePattern module that filters a dataset based on a list of reagents to keep. The module inputs data in a GCT format as well as a list of reagents in a txt file and subsets the data file according to this list.

Parameters

Name	Description
gct file *	File of data to filter in GCT format
keeplist *	list of reagents to keep
extension *	User defined file prefix

^{* -} required

Input Files

- 1. gct file
 - A GCT file containing data to filter.
- 2. keeplist
 - A one-column txt file containing a list of reagents to keep.
- 3. .csv file of known CCLE matches for populating the KnownMatch.CCLE column.

Output Files

1. Filtered GCT data file [prefix]_filtered2.gct

A GCT file containing the data from the input gct file for reagents listed in the keeplist filde.

Platform Dependencies

Task Type:

RNAi

CPU Type:

any

Operating System:

any

Language:

R

Version Comments

Version Release Date Description

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