

# PertCNfilter, v36.2

Filters a gct file of perturbation data based on the copy number of each gene. Uses a per cell line thresholds in an uploaded file.

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**Algorithm Version:** 

#### Introduction

PertCNFilter is a GenePattern module which thresholds data based on copy number thresholds. The user inputs their dataset as a GCT file as well as copy number data as GCT files and threshold information as a text file.

#### **Parameters**

Name	Description
gct file *	data file to filter (.gct)
ABCN file	ABSOLUTE copy number file (.gct)
CN file *	relative copy number (.gct)
threshold file *	Cell line specific thresholds file (.txt)
extension *	filename prefix

<sup>\* -</sup> required

## Input Files

1. gct.file

Data file containing sample information in a GCT file format.

2. ABCN.file

File containing absolute copy number information in a GCT file format.

3. CN.file

GCT file including relative copy number information.

4. threshold.file

Text file describing copy number thresholds for each cell line.

### **Output Files**

1. Copy number filtered data [File prefix]\_CNfiltered.gct Output file after copy number filtering.

## Platform Dependencies

Task Type:

**RNAi** 

**CPU Type:** 

any

**Operating System:** 

any

Language:

R

#### **Version Comments**

Version Release Date Description

