GenePattern

MergeColumns Documentation

Description: Merge datasets by column.
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Summary: Creates a new merged dataset from a list of datasets. Common features (rows) in the datasets are retained while others are discarded.

Parameters:

Name	Description
input filename1	first file to merge - .res, .gct, .snp (non-allele-specific), .cn, .xcn, .odf, or .cls file
input filename2	second file to merge
zip filename	zip file containing files to merge. Either a zip file or the input files above must be provided.
output	name of the merged file

Output Files:
1. File containing the merged datasets. Format of output file is the same as the format of the two input files. If the two input files have different formats, the format of the output file is gct.

Platform dependencies:
      Module type: Preprocess & Utilities
      CPU type: any
      OS: any
      Language: Java