

RXCtoGCT - version 1

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A module for converting tab-delimited text files into the GenePattern .GCT format.

Description:

Reads in a rxc file, counts the number of rows (probes) and columns (samples). Create an output file with standard GCT formatting and row/column counts.

Be sure to check for proper line endings, especially if it looks like there is only one row.

Parameters:

“input”: A tab-delimited text file with L rows and W columns that you want formatted as a GCT.

“output”: The desired output file name. Will have “.gct” appended.