

Principal Components Analysis Documentation

Module name: PCA

Description: Principal Components Analysis

Author: GenePattern Team https://www.genepattern.org/contact

Dana-Farber Cancer Institute

Date: April 28, 2020

Release: 4.2

Performs a principal components analysis. This implementation is an adaptation of the MultiExperiment Viewer PCA analysis (mev.tm4.org).

References:

Raychaudhuri S, Stuart JM, and Altman RB. (2000) Principal components analysis to summarize microarray experiments: application to sporulation time series. Pac. Symp. Biocomput. 455-466.

Parameters:

Name Description

input_filename The input file (.res, .gct, .odf)

output_file The stub for the output files

Return Value:

s matrix
 t matrix
 t matrix
 t matrix
 The matrix of eigenvectors (1 per principal component)

3. u matrix The principal components matrix

4. Stdout.txt: the "stdout" text output from running the program.

Platform dependencies:

Task type: Dimension Reduction

CPU type: any
OS: any
Java JVM level: 1.3
Language: Java