

TCGA.SampleSelection Documentation

Description: Retrieve TCGA data from Broad Firehose and perform sample selection on the basis of expression levels for specific genes of interest for analysis using GSEA tools.

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Summary: Queries cBioPortal for TCGA samples meeting criteria of mRNA expression z-scores relative to all samples (log RNA Seq V2 RSEM) greater than and less than user supplied thresholds. Outputs a GCT file containing TPM (Transcripts per Million) normalized RNA-seq quantifications suitable for ssGSEA, and a CLS file annotating samples as High or Low expression of the gene of interest.

Parameters:

Name	Description
TCGA Collection*	TCGA study cohort (tumor types) to query for sample
	selection.
Gene Symbol*	The HGNC Gene Symbol to use for classifying
	samples as high or low expression
High Expression*	mRNA expression is greater than or equal to this
	threshold for standard deviations above the mean
	will be classified as "high" expression of the selected
	gene.
	(mRNA expression z-scores relative to all samples)
Low Expression*	mRNA expression is less than or equal to this
	threshold for standard deviations below the mean will
	be classified as "low" expression of the selected
	gene.
	(mRNA expression z-scores relative to all samples)
Output Type*	Type of RSEM quantifications to output:
	TPM (transcripts per million, within sample
	normalization, useful for ssGSEA)
	Raw counts (unnormalized counts, usable with
	DESeq2 or other DEG calculations).
MSigDB Version*	MSigDB version to use for Gene Symbol lookup.
	This version should match the version of the gene
	sets intended for all downstream analysis.

^{* =} required

Output File(s): GCT file containing gene expression values in the selected output type for samples that pass the specified thresholds. A CLS file indicating the sample group assignments (high or low expression).

Module Language: R

Source Repository: https://github.com/genepattern/TCGA,SampleSelection/releases/tag/v1

Docker image: bioconductor/bioconductor docker:RELEASE 3 12

Version	Comment
0.1	Initial beta release.