

genefu: a package for breast cancer gene expression analysis

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1 Introduction

The *genefu* package is providing relevant functions for gene expression analysis, especially in breast cancer.

2 Session Info

- R version 2.12.1 (2010-12-16), i686-pc-linux-gnu
- Locale: LC_CTYPE=en_US.utf8, LC_NUMERIC=C, LC_TIME=en_US.utf8, LC_COLLATE=en_US.utf8, LC_MONETARY=C, LC_MESSAGES=en_US.utf8, LC_PAPER=en_US.utf8, LC_NAME=C, LC_ADDRESS=C, LC_TELEPHONE=C, LC_MEASUREMENT=en_US.utf8, LC_IDENTIFICATION=C
- Base packages: base, datasets, graphics, grDevices, methods, stats, tools, utils
- Other packages: cacheSweave 0.4-5, codetools 0.2-8, filehash 2.1-1, formatR 0.2-0, getopt 1.15, highlight 0.2-5, parser 0.0-13, pgfSweave 1.1.3, Rcpp 0.9.2, stashR 0.3-3, tikzDevice 0.5.3
- Loaded via a namespace (and not attached): digest 0.4.2