Package 'msBreast'

February 7, 2011

Type Package
Title Protein Mass Spectra Dataset from a Breast Cancer Study
Version 1.0.3
Date 2011-02-07
Author Lixin Gong, William Constantine, and Yu Alex Chen
Maintainer Lixin Gong <pre><lgong@tibco.com></lgong@tibco.com></pre>
Depends R (>= $2.12.0$)
DependsSplus RSQLite (>= 0.5-6)
Suggests msProcess
Description This package provides a dataset of 96 protein mass spectra generated from a pooled sample of nipple aspirate fluid (NAF) from healthy breasts and breasts with cancer.
License GNU General Public License Version 2
Copyright 2007-2009 TIBCO Software Inc. All rights reserved.
<pre>URL http://www.insightful.com/services/research/proteome/default.asp</pre>
R topics documented:
Breast2003QC 2 Breast2003QCList 3
Index 5

2 Breast2003QC

Breast2003QC

Mass Spectra from a Breast Cancer Quality Control Sample

Description

A data object of class msSet, consisting of 96 mass spectra generated from a pooled sample of nipple aspirate fluid (NAF) from healthy breasts and breasts with cancer. The data set was derived from Breast2003QCList by eliminating the mass region below 950 Da/charge. Please see the references for more details.

Usage

Breast2003QC

Format

An object of msSet. See msSet in package msProcess for details.

Source

http://bioinformatics.mdanderson.org/pubdata.html

References

Coombes, K.R., Tsavachidis, S., Morris, J.S., Baggerly, K.A., and Kuerer, H.M., "Improved peak detection and quantification of mass spectrometry data acquired from surface-enhanced laser desorption and ionization by denoising spectra with the undecimated discrete wavelet transform," *Proteomics*, 5:4107–17, 2005.

Pawlik, T.M., Fritsche, H., Coombes, K.R., Xiao, L., Krishnamurthy, S., Hunt, K.K., Pusztai, L., Chen, J.N., Clarke, C.H., Arun, B., Hung, M.C., and Kuerer, H.M., "Significant differences in nipple aspirate fluid protein expression between healthy women and those with breast cancer demonstrated by time-of-flight mass spectrometry," *Breast Cancer Research and Treatment*, 89(2):149–57, 2005.

Kuerer, H.M., Coombes, K.R., Chen, J.N., Xiao, L., Clarke, C., Fritsche, H., Krishnamurthy, S., Marcy, S., Hung, M.C., and Hunt, K.K., "Association between ductal fluid proteomic expression profiles and the presence of lymph node metastases in women with breast cancer," *Surgery*, 136(5):1061–9, 2004.

Coombes, K.R., Fritsche, Jr., H.A., Clarke, C., Chen, J.N., Baggerly, K.A., Morris, J.S., Xiao, L.C., Hung, M.C., and Kuerer, H.M., "Quality control and peak finding for proteomics data collected from nipple aspirate fluid by surface-enhanced laser desorption and ionization," *Clinical Chemistry*, 49(10):1615–23, 2003.

See Also

Breast2003QCList.

Breast2003QCList 3

Examples

```
if (require("msProcess")) {
  if (!exists("Breast2003QC")) data("Breast2003QC", package="msBreast")

## plot a few spectra
  plot(Breast2003QC, subset=1:5)

## image of all spectra
  image(Breast2003QC)
}
```

Breast2003QCList

Mass Spectra from a Breast Cancer Quality Control Sample

Description

A data object of class msList, consisting of 96 mass spectra generated from a pooled sample of nipple aspirate fluid (NAF) from healthy breasts and breasts with cancer. (see references for details).

Usage

Breast2003QCList

Format

An object of msList. See msList in package msProcess for details.

Source

http://bioinformatics.mdanderson.org/pubdata.html

References

Coombes, K.R., Tsavachidis, S., Morris, J.S., Baggerly, K.A., and Kuerer, H.M., "Improved peak detection and quantification of mass spectrometry data acquired from surface-enhanced laser desorption and ionization by denoising spectra with the undecimated discrete wavelet transform," *Proteomics*, 5:4107–17, 2005.

Pawlik, T.M., Fritsche, H., Coombes, K.R., Xiao, L., Krishnamurthy, S., Hunt, K.K., Pusztai, L., Chen, J.N., Clarke, C.H., Arun, B., Hung, M.C., and Kuerer, H.M., "Significant differences in nipple aspirate fluid protein expression between healthy women and those with breast cancer demonstrated by time-of-flight mass spectrometry," *Breast Cancer Research and Treatment*, 89(2):149–57, 2005.

Kuerer, H.M., Coombes, K.R., Chen, J.N., Xiao, L., Clarke, C., Fritsche, H., Krishnamurthy, S., Marcy, S., Hung, M.C., and Hunt, K.K., "Association between ductal fluid proteomic expression profiles and the presence of lymph node metastases in women with breast cancer," *Surgery*, 136(5):1061–9, 2004.

Coombes, K.R., Fritsche, Jr., H.A., Clarke, C., Chen, J.N., Baggerly, K.A., Morris, J.S., Xiao, L.C., Hung, M.C., and Kuerer, H.M., "Quality control and peak finding for proteomics data collected from nipple aspirate fluid by surface-enhanced laser desorption and ionization," *Clinical Chemistry*, 49(10):1615–23, 2003.

Breast2003QCList

See Also

Breast2003QC.

Examples

```
if (require("msProcess")) {
  if (!exists("Breast2003QCList")) data("Breast2003QCList", package="msBreast")

## print an msList object
Breast2003QCList

## print the synopsis of an msList object
summary(Breast2003QCList)

## plot the 10th spectrum from an msList object
plot(Breast2003QCList, index=10)
```

Index

*Topic datasets Breast2003QC, 2 Breast2003QCList, 3 Breast2003QC, 2, 4 Breast2003QCList, 2, 3 msList, 3 msProcess, 2, 3 msSet, 2