Package 'ClassPrediction'

July 20, 2022

Version 3.1.1

Date 2016-05-09

Title Classes and Methods for ``Class Prediction'' with Microarrays or Proteomics

Author Kevin R. Coombes

Maintainer Kevin R. Coombes <krc@silicovore.com>

Depends R (>= 3.0), GenAlgo (>= 1.0), CrossValidate (>= 2.1)

Suggests Biobase

Description Bundles the classes and pacakges used to help with ``class prediction'' problems in the OOMPA project. Class prediction consists of supervised methods to select relevant features and construct predictive models for binary, ordinal, continuous, or survival outcomes.

License Apache License (== 2.0)

biocViews Microarray, Clustering

URL http://oompa.r-forge.r-project.org/

NeedsCompilation no

R topics documented:

ClassPrediction-package

The ClassPrediction Bundle

Description

This class provides tools to help with class prediction. In its current form, it is a shell used to bundle various other packages that actually do all the work.

Details

Package:	ClassPrediction
Type:	Package
Version:	3.1.1
Date:	2016-05-09
License:	Apache 2.0
LazyLoad:	yes

Author(s)

Kevin R. Coombes <kcoombes@mdanderson.org>

See Also

GenAlg-class, CrossValidate-class.

Index