

Package ‘RvtkStatismo’

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Type Package

Title Integrates statismo and R using the vtkStandardMeshRepresenter

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Description Integrates statismo and R using the vtkStandardMeshRepresenter.
Statismo shape models will be stored as objects of class ``pPCA". (this is work in progress).

License GPL (>=2)

Imports Rcpp (>= 0.11.1),Morpho,Rvcg,methods

LinkingTo Rcpp,RcppEigen

SystemRequirement VTK5.8, statismo (>= 0.9 best ist freshly from github)

URL <http://github.com/zarquon42b/RvtkStatismo>, URL:
<http://github.com/statismo/statismo>

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RvtkStatismo-package *Integrates statismo and R using the vtkStandardMeshRepresente*

Description

Integrates statismo and R using the vtkStandardMeshRepresenter. Statismo shape models will be stored as objects of class "pPCA". (this is work in progress).

Details

Package:	RvtkStatismo
Type:	Package
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Date:	2014-07-01
License:	GPL
LazyLoad:	yes

Author(s)

Stefan Schlager

Maintainer: Stefan Schlager <zarquon42@gmail.com>

References

To be announced

align2domain	<i>align a sample to a model</i>
--------------	----------------------------------

Description

align a sample to a model

Usage

```
align2domain(model, sample, scale = TRUE, ptDomain = NULL,
             ptSample = NULL)
```

```
## S4 method for signature 'pPCA,matrix'
align2domain(model, sample, scale = TRUE,
             ptDomain = NULL, ptSample = NULL)
```

```
## S4 method for signature 'pPCA,mesh3d'
align2domain(model, sample, scale = TRUE,
             ptDomain = NULL, ptSample = NULL)
```

Arguments

model	statistical model of class "pPCA"
sample	matrix or mesh3d
scale	logical: request scaling during alignment
ptDomain	integer vector: specifies the indices of the domain points that are to be used for registration (order is important).
ptSample	integer vector: specifies the indices of the sample that are to be used for registration (order is important).

Value

a rotated (and scaled) mesh or matrix - depending on the input.

ComputeConstrainedModel

Constrains a model of class pPCA by a subset of coordinates

Description

Constrains a model of class pPCA by a subset of coordinates

Usage

```
ComputeConstrainedModel(x, model, align = FALSE, use.lm, deselect = FALSE,
  origSpace = FALSE)
```

Arguments

<code>x</code>	a $k \times 3$ matrix containing the sample's coordinates of the reduced model
<code>model</code>	an object of class <code>pPCA</code>
<code>align</code>	logical: if TRUE, <code>x</code> will be aligned to the models mean
<code>use.lm</code>	integer vector, specifying which coordinates from the full model are to be used/missing (see note)
<code>deselect</code>	logical: if TRUE, <code>use.lm</code> specifies the missing coordinates instead of those present.
<code>origSpace</code>	logical: if <code>align=TRUE</code> and <code>origSpace=TRUE</code> , the representer of the returned model will contain the estimated full shape in the original coordinate system of <code>x</code>

Value

an object of class `pPCA` constrained to `x`

Note

if `deselect = FALSE`, the order of the entries in `use.lm` is interpreted as follows: the i -th entry in `use.lm` specifies the index of the meanshapes coordinate belonging to the i -th coordinate of `x`. if `deselect = TRUE`, the i -th coordinate of `x` is linked to the i -th coordinate of the model's mean with `use.lm` removed.

Examples

```
## create a model superimposed with missing landmarks 3 and 4
require(Morpho)
data(boneData)
newmod <- pPCA(boneLM[,,-1],sigma=0,scale=TRUE,use.lm = 3:4,deselect=TRUE)
## predict the left out shape from the constrained model
boneLM1 <- ComputeConstrainedModel(boneLM[-c(3:4),,1],newmod,align=TRUE,use.lm=3:4,deselect=TRUE,origSpace=TRUE)
## the coordinates of the estimated complete config are now stored in the representer's vertices
## Not run:
##visualize prediction error
deformGrid3d(vert2points(boneLM1$representer),boneLM[, ,1],ngrid=0)

## End(Not run)
```

getCoordVar	<i>get per coordinate variance from a statistical model</i>
-------------	---

Description

get per coordinate variance from a statistical model

Usage

```
getCoordVar(model)
```

Arguments

model object of class pPCA

Note

calculates the per-coordinate variance as described in Luethi(2009)

References

Lüthi M, Albrecht T, Vetter T. 2009. Probabilistic modeling and visualization of the flexibility in morphable models. In: Mathematics of Surfaces XIII. Springer. p 251-264

getDataLikelihood	<i>calculate probability/coefficients for a matrix/mesh given a statistical model</i>
-------------------	---

Description

calculate probability for a matrix/mesh given a statistical model

Usage

```
getDataLikelihood(x, model, align = FALSE, use.lm)
```

```
## S3 method for class 'matrix'  
getDataLikelihood(x, model, align = FALSE, use.lm = NULL)
```

```
## S3 method for class 'mesh3d'  
getDataLikelihood(x, model, align = FALSE, use.lm = NULL)
```

```
getCoefficients(x, model, align = TRUE, use.lm = NULL)
```

Arguments

<code>x</code>	matrix or mesh3d
<code>model</code>	a model of class pPCA
<code>align</code>	logical: if TRUE the data will be aligned to the model's mean
<code>use.lm</code>	integer vector specifying row indices of the coordinates to use for rigid registration on the model's meanshape.

Details

`getDataLikelihood` estimates the likelihood of a dataset for belonging to the model by exploiting the χ^2 -distribution of the (squared) Mahalanobisdistance, which, in turn, is simply the squared norm of the sample's coefficients in the latent space.

Value

`getDataLikelihood` returns a probability, while `getCoefficients` returns the (scaled) scores in the pPCA space.

<code>mesh2vtp</code>	<i>exports a triangular mesh of class mesh3d to a vtp file</i>
-----------------------	--

Description

exports a triangular mesh of class mesh3d to a vtp file

Usage

```
mesh2vtp(mesh, filename = dataname)
```

Arguments

<code>mesh</code>	mesh of class mesh3d
<code>filename</code>	character

meshalign *align meshes stored in a list by their vertices*

Description

align meshes stored in a list by their vertices

Usage

```
meshalign(meshlist, scale = TRUE, use.lm = NULL, deselect = FALSE,
          array = FALSE)
```

Arguments

meshlist	list containing triangular meshes of class "mesh3d"
scale	logical: request scaling during alignment
deselect	logical: if TRUE, missingIndex references the existing coordinates instead of the missing ones.
use.lm	integer vector: specifies the indices of the points that are to be used in the constrained model
array	logical: if TRUE the superimposed vertices will be returned as 3D array.

Value

returns a list of aligned meshes or an array of dimensions $k \times 3 \times n$, where k =number of vertices and n =sample size.

meshlist2array *convert meshes to array consisting of vertex coordinates*

Description

convert meshes to array consisting of vertex coordinates

Usage

```
meshlist2array(meshlist)
```

Arguments

meshlist	list containing triangular meshes of class "mesh3d"
----------	---

Value

returns an array with $k \times 3 \times n$ dimensions where k =number of vertices, and n =sample size.

modelinfo-class *Documentation of class modelinfo*

Description

Documentation of class modelinfo

Usage

```
AddModelInfoParams(x, value)
```

```
## S4 method for signature 'modelinfo'
AddModelInfoParams(x, value)
```

```
SetModelInfoParams(x) <- value
```

Arguments

x an object of class "modelinfo"
value a list of or a single 2-valued character vectors

Details

The class has the following slots

datainfo a list containing 2-valued character vectors

parminfo a list containing 2-valued character vectors

These can be modified using addParams and setParaminfo with

pPCA *calculate or modify a probabilistic PCA based on 3D-coordinates*

Description

calculate or modify a probabilistic PCA based on 3D-coordinates

Usage

```
pPCA(array, align = TRUE, use.lm = NULL, deselect = FALSE, sigma = NULL,
      exVar = 1, scale = TRUE, representer = NULL)
```

```
UpdateModel(model, sigma = NULL, exVar = 1)
```

```
## S4 method for signature 'pPCA'
UpdateModel(model, sigma = NULL, exVar = 1)
```


Arguments

array	array of dimensions $k \times 3 \times n$, where k =number of coordinates and n =sample size.
align	logical: if TRUE, the data will be aligned first
use.lm	integer vector: specifies the indices of the points that are to be used in the constrained model
deselect	logical: if TRUE, use.lm references the missing coordinates instead of the present ones.
sigma	estimate of error variance (sensible is a value estimating coordinate error in terms of observer error)
exVar	numeric value with $0 < \text{exVar} \leq 1$ specifying the PCs to be included by their cumulative explained Variance
scale	logical: allow scaling in Procrustes fitting
representer	a triangular mesh, where the vertices correspond to the coordinates in array, leave NULL for pointclouds.
model	object of class pPCA

Value

returns a probabilistic PCA model as S4 class "pPCA" (see [pPCA-class](#)). UpdateModel is used to modify existing models by changing sigma and exVar.

References

Lüthi M, Albrecht T, Vetter T. 2009. Probabilistic modeling and visualization of the flexibility in morphable models. In: Mathematics of Surfaces XIII. Springer. p 251-264

Examples

```
require(Morpho)
data(boneData)
model <- pPCA(boneLM[, , ])
## change parameters without recomputing Procrustes fit
model1 <- UpdateModel(model, sigma=1, exVar=0.8)
```

Description

Documentation of class pPCA

Details

The class contains the the following slots (still not yet set in stone)

PCA a list containing

- **sdev**: the square roots of the covariance matrix' eigenvalues
- **rotation**: matrix containing the orthonormal PCBasis vectors
- **x**: the scores within the latent space(scaled by 1/sdev)
- **center**: a vector of the mean shape in with coordinates ordered (x1,y1,z1, x2, y2,z2, ..., xn,yn,zn)

scale logical: indicating if the data was aligned including scaling

representer an object of class mesh3d or a list with entry vb being a matrix with the columns containing coordinates and it a 0x0 matrix

sigma the noise estimation of the data

Variance a data.frame containing the Variance, cumulative Variance and Variance explained by each Principal component

rawdata optional data: a matrix with rows containing the mean centred coordinates in order (x1,y1,z1, x2, y2,z2, ..., xn,yn,zn)

PredictSample

predict or restrict a mesh or matrix based on a statistical model

Description

predict or restrict a mesh or matrix based on a statistical model

Usage

```
PredictSample(model, dataset, representer = TRUE, ...)
```

```
## S4 method for signature 'pPCA,matrix,ANY'
```

```
PredictSample(model, dataset, representer = TRUE,
  origSpace = TRUE, use.lm = NULL, deselect = FALSE, sdmax,
  mahaprob = c("none", "chisq", "dist"), align = TRUE, ...)
```

```
## S4 method for signature 'pPCA,mesh3d,logical'
```

```
PredictSample(model, dataset,
  representer = TRUE, origSpace = TRUE, use.lm = NULL, deselect = FALSE,
  sdmax, mahaprob = c("none", "chisq", "dist"), align = TRUE, ...)
```

Arguments

model	model of class pPCA
dataset	a matrix or a mesh3d
representer	if TRUE and the model contains a representer mesh, a surface mesh will be returned, coordinate matrix otherwise.
origSpace	logical: rotate the estimation back into the original coordinate system.
use.lm	optional: integer vector specifying row indices of the coordinates to use for rigid registration on the model's meanshape.
deselect	logical: if TRUE, all BUT the coordinates specified by use.lm will be used for alignment.
sdmax	maximum allowed standard deviation (per Principal axis) within the model space. Defines the probabilistic boundaries.
mahaprob	character: if != "none", use mahalanobis-distance to determine overall probability (of the shape projected into the model space."chisq" uses the Chi-Square distribution of the squared Mahalanobisdistance, while "dist" restricts the values to be within a multi-dimensional sphere of radius sdmax. If FALSE the probability will be determined per PC separately.
align	if TRUE, the sample will be aligned to the mean.
...	currently not in use.

Value

PredictSample returns a matrix/mesh3d restricted to the boundaries given by the modelspace.

See Also

[StatismoModelMembers](#)

read.vtk	<i>imports vtk and vtp files</i>
----------	----------------------------------

Description

imports vtk and vtp files

Usage

```
read.vtk(filename)
```

Arguments

filename	character string
----------	------------------

Value

list of class mesh3d

representer2sample *get the representer from a model of class "pPCA"*

Description

get the representer from a model of class "pPCA"

Usage

```
representer2sample(model)

## S4 method for signature 'pPCA'
representer2sample(model)
```

Arguments

model object of class `pPCA`

Value

an object of class `mesh3d` or `matrix`, depending whether a point cloud or a triangular mesh is the model's representer.

rigidAlign *Fast Procrustes align of coordinates*

Description

Fast Procrustes align of coordinates

Usage

```
rigidAlign(array, scale = TRUE, use.lm = NULL, deselect = FALSE)
```

Arguments

array	array of coordinates
scale	logical: request scaling during alignment
use.lm	integer vector: specifies the indices of the points that are to be used in the constrained model
deselect	logical: if TRUE, use.lm references the missing coordinates instead of the present ones.

Value

a list containing

rotated	array containing registered coordinates
mshape	matrix containing meanshape

Set-pPCA-class *Low level methods to set pPCA class content*

Description

Low level methods to set pPCA class content - not to be invoked directly

Usage

```
SetNoiseVariance(x) <- value
SetRawdata(x) <- value
SetPCBasisMatrix(x) <- value
SetPCsdev(x) <- value
SetMeanVector(x) <- value
SetScores(x) <- value
SetScale(x) <- value
SetPCA(x) <- value
SetVariance(x) <- value

## S4 method for signature 'pPCA'
AddModelInfoParams(x, value)
```

Arguments

model	of class "pPCA"
value	set the specific value
x	an object of class "modelinfo"

Value

returns an updated pPCA object

statismoBuildModel	<i>generate a statistical model using an array of superimposed landmarks or a list of meshes</i>
--------------------	--

Description

generate a statistical model using an array of superimposed landmarks

Usage

```
statismoBuildModel(x, representer, sigma = 0, scale = TRUE)
```

Arguments

x	array of aligned 3D-coordinates or a list of aligned registered meshes.
representer	matrix or triangular mesh of class "mesh3d" with vertices corresponding to rows in the array.
sigma	noise in the data
scale	logical: set to TRUE, if scaling was involved in the registration.

Value

an object of class pPCA ([pPCA-class](#))

See Also

[pPCA](#), [pPCA-class](#), [rigidAlign](#), [meshalign](#)

Examples

```
require(Morpho)
data(boneData)
align <- rigidAlign(boneLM)$rotated
mymod <- statismoBuildModel(align, representer=align[, , 1], sigma=2, scale=TRUE)
## save it
statismoSaveModel(mymod, "mymod.h5")
```

statismoGPmodel	<i>expands a models variability by adding a Gaussian kernel function</i>
-----------------	--

Description

expands a models variability by adding a Gaussian kernel function to the empiric covariance matrix and builds a low-rank approximation of the resulting PCA

Usage

```
statismoGPmodel(model, useEmpiric = TRUE, kernel = list(c(100, 70)),
  ncomp = 10, nystroem = 500)
```

Arguments

model	shape model of class pPCA
useEmpiric	logical: if TRUE, the empiric covariance kernel will be added to the Gaussian ones.
kernel	a list containing two valued vectors containing with the first entry specifying the bandwidth and the second the scaling of the Gaussian kernels.
ncomp	integer: number of PCs to approximate
nystroem	number of samples to compute Nystroem approximation of eigenvectors

Value

returns a shape model of class [pPCA](#)

See Also

[pPCA](#), [pPCA-class](#)

Examples

```
### this is a silly example with only 10 landmarks
require(Morpho)
data(boneData)
align <- rigidAlign(boneLM)$rotated
mod <- statismoBuildModel(align)
GPmod <- statismoGPmodel(mod, kernel=list(c(10,1),c(1,1)))##extend flexibility using two Gaussian kernels
GPmodNoEmp <- statismoGPmodel(mod, kernel=list(c(10,1),c(1,1)), useEmpiric = FALSE)##extend flexibility using two
PC1orig <- DrawSample(mod,2)# get shape in 2sd of first PC of original model
PC1 <- DrawSample(GPmod,2)# get shape in 2sd of first PC of the extended model
PC1NoEmp <- DrawSample(GPmodNoEmp,2)# get shape in 2sd of first PC
##visualize the differences from the mean (green spheres)
deformGrid3d(PC1, DrawMean(GPmod), ngrid=0)##
deformGrid3d(PC1NoEmp, DrawMean(GPmod), ngrid=0, col1=4, add=TRUE)##only deviates in 5 landmarks from the mean (dark
deformGrid3d(PC1orig, DrawMean(GPmod), ngrid=0, col1=5, add=TRUE)
```

statismoLoadModel/statismoSaveModel

save and load a statistical model of class pPCA to statismo hdf5 format

Description

save and load a statistical model of class pPCA to statismo hdf5 format

Usage

```
statismoSaveModel(model, modelName = dataname)
```

```
statismoLoadModel(modelname, scale = FALSE)
```

Arguments

model	object of class pPCA
modelName	filename to read/save
scale	specify if scaling was involved in model generation

Value

statismoLoadModel returns an object of class [pPCA](#) while statismoSaveModel saves an object of class [pPCA](#) to disk in the statismo file format.

See Also

[pPCA](#)

StatismoMatrices

Get Matrices from StatisticalModel class

Description

Get Matrices from StatisticalModel class - such as projection matrices, covariance matrices or Jacobian

Usage

```

GetPCABasisMatrix(model)

GetOrthonormalPCABasisMatrix(model)

GetCovarianceAtPoint(model, pt1, pt2)

GetCovarianceMatrix(model)

GetJacobian(model, pt)

GetProjectionMatrix(model)

## S4 method for signature 'pPCA'
GetPCABasisMatrix(model)

## S4 method for signature 'pPCA'
GetOrthonormalPCABasisMatrix(model)

## S4 method for signature 'pPCA,numeric,numeric'
GetCovarianceAtPoint(model, pt1, pt2)

## S4 method for signature 'pPCA'
GetCovarianceMatrix(model)

## S4 method for signature 'pPCA,numeric'
GetJacobian(model, pt)

## S4 method for signature 'pPCA'
GetProjectionMatrix(model)

```

Arguments

model	model of class "pPCA"
pt	either an integer pointing to the index of the domain or a numeric vector of length 3 specifying a point on the domain of the model
pt1	either an integer pointing to the index of the domain or a numeric vector of length 3 specifying a point on the domain of the model
pt2	either an integer pointing to the index of the domain or a numeric vector of length 3 specifying a point on the domain of the model

Details

see http://statismo.github.io/statismo/classdoc/html/classstatismo_1_1StatisticalModel.html for details.

Value

GetPCABasisMatrix	returns the (scaled) Basis of the latent space
GetOrthonormalPCABasisMatrix	returns the orthonormal Basis of the latent space
GetCovarianceMatrix	returns the covariance matrix - can be huge!!!
GetCovarianceAtPoint	returns the 3 x 3 covariance matrix for pt1 and pt2
GetJacobian	returns the 3 x 3 Jacobian matrix at pt
GetProjectionMatrix	returns matrix to project a sample vector into the latent space (this is not a member function but might prove useful anyway)

StatismoModelMembers *Implementation/Emulation of the statismo StatisticalModel class.*

Description

Implementation/Emulation of the statismo StatisticalModel class.

Usage

```

DrawMean(model)

DrawMeanAtPoint(model, pt)

DrawSample(model, coefficients = NULL, addNoise = FALSE)

DrawSampleVector(model, coefficients, addNoise = FALSE)

DrawSampleAtPoint(model, coefficients, pt, addNoise = FALSE)

ComputeCoefficientsForDataset(model, dataset)

ComputeCoefficientsForPointValues(model, sample, pt, ptNoise = 0)

GetDomainPoints(model)

GetDomainSize(model)

EvaluateSampleAtPoint(model, sample, pt)

GetModelInfo(model)

```

```
## S4 method for signature 'pPCA'  
DrawMean(model)  
  
## S4 method for signature 'pPCA,numeric'  
DrawMeanAtPoint(model, pt)  
  
## S4 method for signature 'pPCA'  
DrawSample(model, coefficients = NULL, addNoise = FALSE)  
  
## S4 method for signature 'pPCA'  
DrawSampleVector(model, coefficients, addNoise = FALSE)  
  
## S4 method for signature 'pPCA,numeric,numeric'  
DrawSampleAtPoint(model, coefficients, pt,  
  addNoise = FALSE)  
  
## S4 method for signature 'pPCA'  
ComputeCoefficientsForDataset(model, dataset)  
  
## S4 method for signature 'pPCA'  
GetDomainPoints(model)  
  
## S4 method for signature 'pPCA'  
GetDomainSize(model)  
  
## S4 method for signature 'pPCA,matrix,numeric,numeric'  
ComputeCoefficientsForPointValues(model,  
  sample, pt, ptNoise = 0)  
  
## S4 method for signature 'pPCA,matrix,matrix,numeric'  
ComputeCoefficientsForPointValues(model,  
  sample, pt, ptNoise = 0)  
  
## S4 method for signature 'pPCA,numeric,numeric,numeric'  
ComputeCoefficientsForPointValues(model,  
  sample, pt, ptNoise = 0)  
  
## S4 method for signature 'pPCA,matrix,numeric'  
EvaluateSampleAtPoint(model, sample, pt)  
  
## S4 method for signature 'pPCA,mesh3d,numeric'  
EvaluateSampleAtPoint(model, sample, pt)  
  
## S4 method for signature 'pPCA'  
GetModelInfo(model)
```

Arguments

model	object of class <code>pPCA</code>
dataset	an (already aligned) mesh or $k \times 3$ matrix containing the datasets coordinates.
coefficients	specify coefficients in the latent space to draw a sample
addNoise	logical: if TRUE noise as specified in the model will be added to the returned sample
ptNoise	specify the noise estimated in the points.
sample	depending on the function a matrix, a numeric vector or a mesh3d (see methods below)
pt	either an integer pointing to a coordinate or a 3D-vector containing the coordinates of the domain point of interest. For <code>ComputeCoefficientsForPointValues</code> , this can also specify a matrix of coordinates on the domain.

Details

see http://statismo.github.io/statismo/classdoc/html/classstatismo_1_1StatisticalModel.html for details.

Value

<code>DrawMean</code>	Get the mean (either a matrix or a mesh3d)
<code>GetMeanVector</code>	Get the mean vector
<code>DrawMeanAtPoint</code>	Get a specific point of the mean (numeric vector)
<code>DrawSample</code>	Draw a sample from the model (either a matrix or a mesh3d)
<code>DrawMeanAtPoint</code>	Get a specific point of the mean (numeric vector)
<code>DrawSampleAtPoint</code>	Draw a sample of a specific point from the model (numeric vector)
<code>ComputeCoefficientsForDataset</code>	Computes the coefficients of the latent variables
<code>ComputeCoefficientsForPointValues</code>	Returns the coefficients of the latent variables for the given values provided in two $k \times 3$ matrices or two vectors of length 3, or one matrix/vector and a vector containing the indices on the domain corresponding to these points
<code>GetDomainPoints</code>	a matrix containing the points of the model's domain
<code>GetDomainSize</code>	get the size of the model's domain
<code>EvaluateSampleAtPoint</code>	Returns the value of the given sample at the point specified (either as point on the domain or as an index)

StatismoParameters *Get model parameters*

Description

Get model parameters such as variance or noise variance

Usage

```
GetNoiseVariance(model)
```

```
GetMeanVector(model)
```

```
GetPCAVarianceVector(model)
```

```
## S4 method for signature 'pPCA'  
GetNoiseVariance(model)
```

```
## S4 method for signature 'pPCA'  
GetMeanVector(model)
```

```
## S4 method for signature 'pPCA'  
GetPCAVarianceVector(model)
```

Arguments

model model of class "pPCA"

Details

see http://statismo.github.io/statismo/classdoc/html/classstatismo_1_1StatisticalModel.html for details.

Value

GetNoiseVariance returns the estimated noise in the model

GetPCAVarianceVector returns the variance in the model

GetMeanVector returns the model's mean vector

statismoReducedVariance

Reduce an existing statistical shape model

Description

Reduce an existing statistical shape model either to first n PCs or by explained Variance

Usage

```
statismoReducedVariance(model, exVar = 1, npc = 0, scores = TRUE)
```

Arguments

model	
exVar	restricts model by explained variance - with $0 < \text{exVar} < 1$
npc	number of PCs retained in the model (overrides exVar)
scores	logical: request recomputation of PC-scores

Examples

```
require(Morpho)
data(boneData)
align <- rigidAlign(boneLM)$rotated
mymod <- statismoBuildModel(align,representer=align[, , 1],sigma=2,scale=TRUE)
reducemod <- statismoReducedVariance(mymod,0.5)
```

StatismoSample

Retrieve information about a sample from the model

Description

Retrieve information about a sample from the model

Usage

```
ComputeLogProbabilityOfDataset(model, dataset)
```

```
ComputeProbabilityOfDataset(model, dataset)
```

```
## S4 method for signature 'pPCA'
ComputeLogProbabilityOfDataset(model, dataset)
```

```
## S4 method for signature 'pPCA'
ComputeProbabilityOfDataset(model, dataset)
```


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