

# Package ‘RvtkStatismo’

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**Type** Package

**Title** Integrates statismo and R using the vtkStandardMeshRepresenter

**Version** 0.2.140701

**Date** 2014-07-01

**Author** Stefan Schlager, the authors of Statismo

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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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**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
Version:	0.2.140701
Date:	2014-07-01
License:	GPL
LazyLoad:	yes

**Author(s)**

Stefan Schlager  
Maintainer: Stefan Schlager <zarquon42@gmail.com>

**References**

To be announced

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align2domain	<i>align a sample to a model</i>
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**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.

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ComputeConstrainedModel	<i>Constrains a model of class pPCA by a subset of coordinates</i>
-------------------------	--

---

**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 meanshape's 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

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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  $\chi^2$ -distribution of the (squared) Mahalanobis distance, 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	<i>align meshes stored in a list by their vertices</i>
-----------	--

---

**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	<i>convert meshes to array consisting of vertex coordinates</i>
----------------	---

---

**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	<i>Documentation of class modelinfo</i>
-----------------	---

---

### Description

Documentation of class modelinfo

### Usage

```
AddModelInfoParams(x, value)

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

SetModelInfoParams(x) <- value
```

### Arguments

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	<i>calculate or modify a probabilistic PCA based on 3D-coordinates</i>
------	--

---

### 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 follwing 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	<i>get the representer from a model of class "pPCA"</i>
--------------------	---

---

### 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 <code>pPCA</code>
-------	-----------------------------------

### Value

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

---

rigidAlign	<i>Fast Procrustes align of coordinates</i>
------------	---

---

### 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	<i>Low level methods to set pPCA class content</i>
----------------	--

---

**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

**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 <a href="#">pPCA</a>
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 <a href="#">pPCA</a>
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](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](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	<i>Get model parameters</i>
--------------------	-----------------------------

---

## 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](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)
```

**Arguments**

model	model of class "pPCA"
dataset	a matrix or mesh3d aligned to the model's mean

**Value**

ComputeLogProbabilityOfDataset	returns the log-probability density for the sample
ComputeProbabilityOfDataset	returns the probability density for the sample

**See Also**

[getDataLikelihood](#)

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