Package ‘LS2Wstat’

September 17, 2023

Type Package
Title A Multiscale Test of Spatial Stationarity for LS2W Processes
Version 2.1-5
Date 2023-08-20
Depends LS2W(>= 1.3-1)
Imports matrixStats, spdep
License GPL-2
Author Sarah Taylor [aut],
    Matt Nunes [aut, cre],
    Idris Eckley [ctb, ths]
Maintainer Matt Nunes <nunesrpackages@gmail.com>
LazyLoad yes
NeedsCompilation no
Repository CRAN
Date/Publication 2023-09-17 08:52:36 UTC

R topics documented:

  LS2Wstat-package . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  2
  avespecvar . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  2
  compareImages . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  3
  countTextures . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  5
  cropimage . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  6
  getpval . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  7
  imageQT . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .  8
  lincurve . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . 10
  mix2images . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . 11
  plot.imageQT . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . 12
  plotmtx . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . 13
Description

This package contains functions for testing for stationarity within images, specifically locally stationary wavelet (LS2W) fields. In addition the package contains functions for implementing quadtree image decompositions, as well as code for simulating LS2W processes for a given spectral structure.

Author(s)

Sarah L. Taylor and Matt Nunes
Maintainer: Matthew Nunes <nunesrpackages@gmail.com>

References

For details on testing LS2W fields for stationarity, see

For further information on LS2W processes, see

avespecvar

A test statistic for spatial stationarity.

Description

Calculates a test statistic for a test of stationarity based on the local wavelet spectrum.

Usage

avespecvar(spectrum)
**compareImages**

Arguments

- **spectrum**: A local wavelet spectrum estimate, i.e. a cddews object.

Details

The test statistic given by Taylor et al. (2014) for a test for stationarity is computed for use in the bootstrap testing procedure (**TOS2D**).

Value

- **statistic**: The value of the test statistic for the given spectrum.

Author(s)

Sarah L. Taylor

References


See Also

- **TOS2D**

Examples

```r
#Generate a cddews object
#
X <- Haar2MA.diag(64)

testspec<-cddews(X,smooth=FALSE)

#Find the value of the test statistic
#
avespecvar(testspec)
#
```

---

**compareImages**

*Assesses whether two textured images are the same texture.*

Description

The function combines two images together, and then tests the montage for stationarity.

Usage

```r
compareImages(Im1, Im2, testsize = min(nrow(Im1), nrow(Im2)), alpha=0.05,...)
```
Arguments

- **Im1**: The first image to be compared.
- **Im2**: The second image to be compared.
- **testsize**: The size of the combined image montage to be tested for stationarity.
- **alpha**: The significance of the stationarity test.
- **...**: Any other optional arguments to `TOS2D`.

Details

An image montage of two images is created, and the homogeneity measure TOS2D is used in combination with `getpval` to assess stationarity of the montage. If the image is assessed as stationary, the two images are considered as the same texture.

Value

- **montageres**: A boolean value indicating whether the montage of Im1 and Im2 is stationary.

Author(s)

Sarah L. Taylor and Matt Nunes

References


See Also

- `TOS2D`, `countTextures`

Examples

```r
# create two images to be compared:
X1<-simTexture(32,K=1,imtype="S1")[[1]]
X2<-simTexture(32,K=1,imtype="S1", sd=1.6)[[1]]

# use the test to compare them:

test<-compareImages(X1,X2,nsamples=100, smooth=FALSE)
```
countTextures

Description

Groups a list of (stationary) images into texture classes.

Usage

countTextures(Imgs, medpol = TRUE, ...)

Arguments

Imgs A list of images to classify into textures.
medpol A boolean value indicating whether to zero mean the images (with Tukey's median polish) prior to classification.
... Any other optional arguments to the classification function compareImages.

Details

The procedure recursively uses the function compareImages to decide whether two images are of the same texture or not. More specifically, the first image is sequentially tested with all others in the list, assigning the images the label "1" if assessed as the same texture as the first image. All other (unclassified) images are then similarly compared with candidates from different texture classes, until all images have been assigned a group label. Testing recursively in this way, there are at most \( \binom{\text{length(Imgs)} \times 2} {2} \) comparisons performed, but in reality the number could be a lot fewer.

Value

Iclass A vector (of length length(Imgs)) of texture labels corresponding to each image in Imgs.

Author(s)

Sarah L. Taylor and Matt Nunes

References


See Also

compareImages
Examples

```r
## Not run:
X1<-simTexture(128,K=1,imtype="S1")[[1]]
X2<-simTexture(128,K=1,imtype="S1")[[1]]
X3<-simTexture(128,K=1,imtype="S1",sd=1.6)[[1]]
Xlist<-list(X1,X2,X3)
Xlist.class<-countTextures(Xlist, bs=100)
## End(Not run)
```

cropimage

Crops a rectangular image to a specified (square) dimension

Description

If the input image is not of dimension $2^n \times 2^n$, for some $n$, then the image is cropped to an optionally specified size.

Usage

```r
cropimage(image, newsize = NULL, pos = "e")
```

Arguments

- `image`: The image you wish to crop.
- `newsize`: An optional dimension (smaller than the original image dimension), to which the image should be cropped.
- `pos`: The position of the subimage to take when cropping an image. See the documentation for `mix2images` for more details.

Details

As we often wish to work with images whose dimensions are some power of 2, this function will determine whether the image is of an appropriate size and if not it will crop the image so that it is. The optional `pos` argument specifies the position of the cropped subimage to be returned; for example `pos="e"` specifies the central region. See `mix2images` for more details on the positioning argument.

Value

- `subim`: A square image with dimension $2^n \times 2^n$, where $n$ is either `newsize` or the largest feasible dyadic power smaller than the original image dimensions.
getpval

Author(s)
Matt Nunes

See Also
mix2images

Examples

# Create an image with dimensions not a power of two
#
testimage <- matrix(rnorm(300^2),nrow=300,ncol=300)
#
# Crop the image
#
Newimage <- cropimage(testimage)
#
# Check new dimension size.
#
dim(Newimage)
#

getpval

Computes a p-value for the output of the test for stationarity.

Description

Computes and returns a p-value from the output of the bootstrap test for stationarity.

Usage

getpval(statvec, verbose = TRUE)

Arguments

statvec A vector of test statistics, such as that given by TOS2D. The first value must be
the value of the test statistic for the original image.

verbose If TRUE then the p-value is printed and a sentence declaring "stationary" or "not
stationary" is printed.

Value

p The p-value of the test.

Author(s)
Sarah L. Taylor
References


See Also

TOS2D

Examples

```r
# Generate a stationary image

testimage <- matrix(rnorm(64*64), nrow=64, ncol=64)

# Run test of stationarity

## Not run: TestofStat<-TOS2D(testimage)

# Obtain p-value

gotpval(TestofStat$samples)

## End(Not run)
```

---

**imageQT**

*Performs an image quadtree decomposition.*

Description

The quadtree decomposition is achieved by recursively splitting subimages into regions of stationarity.

Usage

```r
imageQT(image, test = TOS2D, minsize = 64, alpha = 0.05, ...)
```

Arguments

- **image**: An image to be decomposed.
- **test**: A function for assessing regions of spatial homogeneity, for example TOS2D.
- **minsize**: The smallest region to test for homogeneity.
- **alpha**: The significance level for the homogeneity test `test`.
- **...**: Any other (optional) arguments to TOS2D.
Details

This function works by assessing an image for homogeneity. If it is not homogeneous, the image is split into its four subquadrants. Each of these is then tested for homogeneity. The heterogeneous subimages are then again subdivided and tested again. This procedure is repeated until either all subimages are deemed stationary or the minimum testing size minsize is reached.

Value

An object of class imageQT with the following components:

- **data.name**: The image analysed.
- **indl**: The index representation of the nonstationary images in the quadtree decomposition.
- **resl**: The results of the stationarity testing (from binfun) during the quadtree decomposition. The results giving 0 match those contained in the indl component and the results giving 1 match those contained in the indS component.
- **imsize**: The original image dimension.
- **imS**: The stationary subimages in the quadtree decomposition.
- **indS**: The index representation of the stationary images in the quadtree decomposition.
- **minsize**: The minimum testing region used during the quadtree decomposition.

Author(s)

Sarah L. Taylor and Matt Nunes

References


See Also

`plot.imageQT`, `TOS2D`

Examples

```r
# generate an image:
X<-simTexture(128,K=1,imtype="NS1", sd = 3)[[1]]

# Not run: XQT<-imageQT(X,binfun=TOS2D.bin)
```
**lincurve**

A linear function between two constant values.

---

**Description**

A function with which to generate nonstationary covariance structure.

**Usage**

```r
lincurve(x, start = 1, end = 2, a = 0.25)
```

**Arguments**

- **x**: a sequence of x-values.
- **start**: a starting value for the linear function.
- **end**: an ending value for the linear function.
- **a**: a proportion of the x-values for the linear part of the function.

**Value**

- **y**: the y-values associated to the linear function.

**Author(s)**

Matt Nunes

**See Also**

`scurve, simTexture`

**Examples**

```r
x <- seq(0, 1, length=128)

y <- lincurve(x, start=1, end=2, a=.25)

plot(x, y, type="l")
```
**mix2images**

Insert one image into another.

**Description**
Image A is re-sized to a specified proportion of Image B, then inserted into Image B at a given position.

**Usage**
mix2images(imageA, imageB, prop = 0.25, pos = "e")

**Arguments**
- **imageA**: The first image which is resized and placed inside the second image.
- **imageB**: The second image, into which the first is placed.
- **prop**: The proportion of Image B to be taken up by Image A.
- **pos**: The exact position of image A in image B. Possible options are "a", "b", "c", "d", "e" which corresponds to (a) top-right, (b) bottom-right, (c) top-left, (d) bottom-left and (e) centred. A more exact location may be specified by inputting pos=c(x,y), which represents the position in pixels from the top-left of the image (i.e. c(x,y) puts Image A x pixels down and y pixels across from the top-left corner of Image B.)

**Details**
This function first of all crops Image A to be a given proportion of Image B and then inserts it into image B at the specified location. If image B is too small for the size of image A required then the whole of image A is placed in image B. Both must be dyadic in length and square images.

**Value**
- **ImageB**: A matrix with the specified values selected exchanged to those of Image A.

**Author(s)**
Sarah L. Taylor

**References**
Examples

```r
# Generate 2 images.
#
ImageA <- matrix(rnorm(256^2), nrow=256, ncol=256)
ImageB <- matrix(rnorm(256^2, sd=2.8), nrow=256, ncol=256)
#
# Insert Image A into Image B at a proportion of 0.25
#
MixImaImb <- mix2images(ImageA, ImageB, prop=0.25, pos="e")
#
```

plot.imageQT

A plot function for quadtree decompositions.

Description

A plot function for quadtree decompositions.

Usage

```r
## S3 method for class ‘imageQT’
plot(x, cires, unclassval = 0, class = FALSE, QT = TRUE,
    return = FALSE, qtl = 1, …)
```

Arguments

- `x` A quadtree decomposition object, such as output from `imageQT`.
- `cires` Results of `countTextures` for the classification of subimages produced by the quadtree decomposition.
- `unclassval` A value for unclassified values in a quadtree decomposition.
- `class` A boolean value indicating whether to plot the results from `countTextures`.
- `QT` A boolean value indicating whether to plot the quadtree decomposition.
- `return` A boolean value indicating whether to return the matrix associated to the plotted image.
- `qtl` Colour specification for the lines drawn in the image segmentation (for `QT=TRUE`).
- `…` Any other optional arguments to `image`.

Details

The function plots the chosen quadtree decomposition, and optionally the textured region classification output from `countTextures`. If the classification output is plotted (`class=TRUE`), each textured region is uniquely coloured according to its texture group.

Value

- `immat` the matrix associated to the plotted image.
plotmtx

Author(s)
Sarah L. Taylor and Matt Nunes

See Also
imageQT, countTextures

Examples

## Not run:
X<-simTexture(256,K=1,imtype="NS2")[[1]]
XQT<-imageQT(X, bs=100, smooth=FALSE)
XCI <- Tex(XQT$imS, bs=100, smooth=FALSE)
plot(XQT, XCI, QT=T, class=T)
## End(Not run)

plotmtx  Image manipulation

Description
A function which rearranges image content for nice plotting.

Usage
plotmtx(m)

Arguments
m  An image (matrix) for converting so that it can be plotted.

Details
Due to the input and plotting output of the R base function image, this function reorders the pixels within an image such that, when used, the image function produces a plot of a image (matrix) "as is".

Value
m.out  The manipulated image corresponding to the input image.
Author(s)
Matt Nunes

See Also
image

Examples

Im<-simTexture(n=256,type="NS4",K=1)[[1]]

image(plotmtx(Im))

print.imageQT

Print out information about a imageQT object in readable form.

Description
This function prints out information about a imageQT object in a nice human-readable form.

Usage

## S3 method for class 'imageQT'
print(x, ...)

Arguments

x An object of class 'imageQT' about which you wish to print information.

... This argument actually does nothing in this function!

Author(s)
Matt Nunes

See Also
imageQT
Examples

## Not run:
#
# Generate a imageQT object for a HaarMontage realisation
#
X<-simTexture(n=256,K=1,imtype="S1")[[1]]

Xres <- imageQT(X)

print(Xres)

## End(Not run)

print.TOS2D

Print out information about a TOS2D object in readable form.

Description

This function prints out information about a TOS2D object in a nice human-readable form.

Usage

## S3 method for class 'TOS2D'
print(x, ...)

## S3 method for class 'TOS2D'
summary(object, ...)

Arguments

x, object
An object of class 'TOS2D' about which you wish to print information.

... This argument actually does nothing in this function!

Author(s)

Matt Nunes

See Also

TOS2D

Examples

## Not run:
#
# Generate a TOS2D object for a HaarMontage realisation
#
X<-simTexture(n=256,K=1,imtype="S1")[[1]]
scurve <- TOS2D(X)
summary(Xres)
## End(Not run)

---

**scurve**  
*An S curve function between two constant values.*

**Description**  
A function with which to generate nonstationary covariance structure.

**Usage**  
scurve(x, a = 1, start = 1, end = 2)

**Arguments**  
- **x**: a sequence of x-values.  
- **a**: The coefficient of slope of the curve.  
- **start**: a starting value for the curve  
- **end**: an ending value for the curve

**Value**  
y: the function values associated to x depicting an S-curve.

**Author(s)**  
Matt Nunes

**See Also**  
lincurve, simTexture

**Examples**  
x <- seq(0, 1, length = 100)

```r
y <- scurve(x, .4, 1, 2)
plot(x, y, type = "l")
```
**simTexture**

*simTexture*ulation function for LS2W processes.

**Description**

This function will generate images of a specified type

**Usage**

```r
simTexture(n = 256, sd = 1, K = 150, imtype = "S1", ...)
```

**Arguments**

- `n` The dimension of the image to be generated.
- `sd` The standard deviation of the increments of the LS2W process to be generated.
- `K` The number of images to generate.
- `imtype` The type of image(s) to create. Must be one of "S1", "S2", "S3", "S4", "NS1", "NS2", "NS3", "NS4", "NS5", "NS6", "NS7". See details for descriptions of the processes.
- `...` Any other optional arguments needed for the image generation (see details).

**Details**

Several different processes can be generated with the `simTexture` function. The stationary processes are: a random normal process of specified standard deviation, `sd` (S1); a spatial moving average process with parameter `rho` (S2); an isotropic random field with a Matern covariance with shape parameter `nu` (S3) and a diagonal Haar moving average process of a specified order `order` and standard deviation `sd` (S4) (see the `Haar2MA.diag` function in the LS2W package for more details).

The nonstationary processes are: a random field with unit standard deviation on the first half-plane, concatenated with a random normal half-plane of standard deviation `sd` (NS1); a white noise half-plane concatenated with a Matern stationary process (NS2); a *Haar Montage* of specified standard deviation `sd` (NS3) (see the LS2W `HaarMontage` function for more details); a process with a slowly-varying covariance structure (NS4); a white noise process with a central subregion of random Normal deviates with non-unit standard deviation `sd` (NS5); a white noise process with a subregion of random Normal deviates with non-unit standard deviation in the middle section of the top left quadrant `sd` (NS6); the final process is similar to NS5, except that there is an additional texture in a subregion of the image. In other words, the image is a montage of three two-dimensional Normal processes with differing standard deviations. The base texture is again of unit variance, whereas the other two textures have standard deviations `sd` and `sd2` (NS7).

The other optional arguments for `simTexture` are as follows:
- `type` - the type of neighbourhood dependence for the random field, either "queen" or "rook" (see the `cell2nb` function documentation in the spdep package for more details).
- `rho` - moving average parameter for the process S2.
- `nu` - shape parameter for the Matern covariance for process S3.
- `order` - Haar moving average order for S4.
fn - s curve or lincurve for NS4.
start - start value for NS4 (passed into s curve or lincurve).
end - end value for NS4 (passed into s curve or lincurve).
a - "gradient" for NS4 (passed into s curve or lincurve).
prop - proportion of inserted subimage for NS5, NS6 and the first subimage (NS7).
sd2 - standard deviation of second inserted subimage for NS7.
prop2 - proportion of second inserted subimage for NS7.
pos1 - position of first inserted subimage for NS7.
pos2 - position of second inserted subimage for NS7.

Value

images A list of length K, with each list entry being an image of dimension n x n with the chosen spectral structure.

Warning

Generating lots of images of high dimension may take a long time!
Note that as of version 2.1-4 (2022-05-09), textures S3 and NS2 have been temporarily removed (commented out) from the functionality of this function, due to a broken package (RandomFields).

Author(s)

Sarah L. Taylor and Matt Nunes

References

Matern, B. (1960) Spatial variation. Stochastic models and their application to some problems in forest surveys and other sampling investigations Meddeleanden fran statens Skogsforskningsinstitut 49 (5).

See Also

HaarMontage

Examples

X1 <- simTexture(128,K=1,imtype="S4",order=3)
X2 <- simTexture(128,K=1,imtype="NS4",fn=lincurve,a=.25,start=1,end=2)
X3 <- simTexture(128,K=1,imtype="NS5",sd=1.6,prop=.25)
X4 <- simTexture(128,K=1,imtype="NS7",sd=1.6,prop=.25, sd2=2.8, prop2=0.25, pos1=c(10,10),pos2="e")

# try plotting the images:
## Not run: image(plotmtx(X1[[1]]))

**TOS2D**

Perform bootstrap stationarity test for images.

**Description**

For a given image this function performs bootstrapping to test the hypothesis that the image is stationary.

**Usage**

```r
TOS2D(image, detrend = FALSE, nsamples = 100, theTS = avespecvar, verbose = TRUE,...)
```

**Arguments**

- `image`: The image you want to analyse.
- `detrend`: This specifies whether to use Tukey’s median polish to remove the image trend.
- `nsamples`: Number of bootstrap simulations to carry out.
- `theTS`: Specifies the particular test statistic to be used. This function should measure the departure from constancy of the wavelet spectrum.
- `verbose`: If TRUE informative messages are printed.
- `...`: Any other arguments supplied to the LS2W function `cddews`.

**Details**

This function first of all crops the image (if necessary) to have dyadic dimensions. The test statistic (theTS), which should be based upon the local wavelet spectrum, is calculated for this original image and the local wavelet spectrum under the null hypothesis is calculated, so as to be able to simulate realisations under the null hypothesis. `nsamples` images are simulated and test statistic is found for each. The function returns all the test statistic values which may be passed to `getpval` in order to find a p-value for the test. For full details on this testing procedure see Taylor et al. (2014).

**Value**

A list with the following components:

- `data.name`: The name of the image analysed.
- `samples`: A vector of length `nsamples+1`. The first entry is the value of the test statistic computed on the original image while the remaining entries are test statistic values for the simulated images.
- `statistic`: The name of the test statistic used.
- `p.value`: The bootstrap p-value for the test.
Author(s)
Sarah L. Taylor

References

See Also
avespecvar, getpval

Examples

# Generate a stationary image
#
testimage <- matrix(rnorm(64*64), nrow=64, ncol=64)
#
#Run test of stationarity

## Not run: TestofStat<TOS2D(testimage)
Index

* datagen
  simTexture, 17
* manip
  compareImages, 3
cropImage, 6
imageQT, 8
lincurve, 10
mix2images, 11
plotmtx, 13
scurve, 16
* methods
  countTextures, 5
* nonparametric
  avespecvar, 2
* package
  LS2Wstat-package, 2
* plot
  plot.imageQT, 12
* print
  print.imageQT, 14
print.TOS2D, 15
* statistic
  getpval, 7
TOS2D, 19
avespecvar, 2, 20

cell2nb, 17
compareImages, 3, 5
countTextures, 4, 5, 13
cropImage, 6
getpval, 7, 19, 20

Haar2MA.diag, 17
HaarMontage, 17, 18

image, 12–14
imageQT, 8, 13, 14
lincurve, 10, 16

LS2Wstat (LS2Wstat-package), 2
LS2Wstat-package, 2

mix2images, 6, 7, 11

plot.imageQT, 9, 12
plotmtx, 13
print.imageQT, 14
print.TOS2D, 15

scurve, 10, 16
simTexture, 10, 16, 17
summary.TOS2D (print.TOS2D), 15

TOS2D, 3, 4, 7–9, 15, 19

21