

Package ‘aws’

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Title Adaptive Weights Smoothing

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Imports methods, gsl

Description We provide a collection of R-functions implementing adaptive smoothing procedures in 1D, 2D and 3D. This includes the Propagation-Separation Approach to adaptive smoothing, the Intersecting Confidence Intervals (ICI), variational approaches and a non-local means filter. The package is described in detail in Polzehl J, Papafitsoros K, Tabelow K (2020). Patch-Wise Adaptive Weights Smoothing in R. Journal of Statistical Software, 95(6), 1-27. <doi:10.18637/jss.v095.i06>, Usage of the package in MR imaging is illustrated in Polzehl and Tabelow (2023), Magnetic Resonance Brain Imaging, 2nd Ed. Appendix A, Springer, Use R! Series. <doi:10.1007/978-3-031-38949-8>.

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aws-package

Adaptive Weights Smoothing

Description

We provide a collection of R-functions implementing adaptive smoothing procedures in 1D, 2D and 3D. This includes the Propagation-Separation Approach to adaptive smoothing, the Intersecting Confidence Intervals (ICI), variational approaches and a non-local means filter. The package is described in detail in Polzehl J, Papafitsoros K, Tabelow K (2020). Patch-Wise Adaptive Weights Smoothing in R. Journal of Statistical Software, 95(6), 1-27. <doi:10.18637/jss.v095.i06>, Usage of the package in MR imaging is illustrated in Polzehl and Tabelow (2023), Magnetic Resonance Brain Imaging, 2nd Ed. Appendix A, Springer, Use R! Series. <doi:10.1007/978-3-031-38949-8>.

Details

The DESCRIPTION file:

```
Package:      aws
Version:      2.5-7
Date:         2026-07-13
Title:        Adaptive Weights Smoothing
Authors@R:    c(person("Joerg","Polzehl",role=c("aut","cre"),email="joerg.polzehl@wias-berlin.de"),person("Felix","Anker","Anker",role="cre"),person("Joerg","Polzehl",role="aut"),person("Felix","Anker",role="aut"))
Author:       Joerg Polzehl [aut, cre], Felix Anker [ctb]
Maintainer:   Joerg Polzehl <joerg.polzehl@wias-berlin.de>
Depends:      R (>= 3.4.0), awsMethods (>= 1.1-1)
Imports:      methods, gsl
Description:   We provide a collection of R-functions implementing adaptive smoothing procedures in 1D, 2D and 3D. This package is Copyright (C) 2005-2024 Weierstrass Institute for Applied Analysis and Stochastics.
License:      GPL (>=2)
Copyright:    This package is Copyright (C) 2005-2024 Weierstrass Institute for Applied Analysis and Stochastics.
URL:          https://www.wias-berlin.de/people/polzehl/
RoxygenNote:  5.0.1
```

Index of help topics:

ICIcombined	Adaptive smoothing by Intersection of Confidence Intervals (ICI) using multiple windows
ICIsmooth	Adaptive smoothing by Intersection of Confidence Intervals (ICI)
ICIsmooth-class	Class '"ICIsmooth"'
TV_denoising	TV/TGV denoising of image data
aws	AWS for local constant models on a grid
aws-class	Class '"aws"'
aws-package	Adaptive Weights Smoothing
aws.gaussian	Adaptive weights smoothing for Gaussian data with variance depending on the mean.
aws.irreg	local constant AWS for irregular (1D/2D) design
aws.segment	Segmentation by adaptive weights for Gaussian models.
awsLocalSigma	3D variance estimation
awsdata	Extract information from an object of class aws
awssegment-class	Class '"awssegment"'
awstestprop	Propagation condition for adaptive weights smoothing
awsweights	Generate weight scheme that would be used in an additional aws step
binning	Binning in 1D, 2D or 3D
extract-methods	Methods for Function 'extract' in Package 'aws'
gethani	Auxiliary functions (for internal use)
kernsm	Kernel smoothing on a 1D, 2D or 3D grid
kernsm-class	Class '"kernsm"'

lpaws	Local polynomial smoothing by AWS
nlmeans	NLMeans filter in 1D/2D/3D
paws	Adaptive weights smoothing using patches
plot-methods	Methods for Function 'plot' from package 'graphics' in Package 'aws'
print-methods	Methods for Function 'print' from package 'base' in Package 'aws'
qmeasures	Quality assessment for image reconstructions.
risk-methods	Compute risks characterizing the quality of smoothing results
show-methods	Methods for Function 'show' in Package 'aws'
smooth3D	Auxiliary 3D smoothing routines
smse3ms	Adaptive smoothing in orientation space SE(3)
summary-methods	Methods for Function 'summary' from package 'base' in Package 'aws'
vaws	vector valued version of function 'aws' The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient likelihood models with vector valued response on a 1D, 2D or 3D grid.
vpaws	vector valued version of function 'paws' with homogeneous covariance structure

Author(s)

Joerg Polzehl [aut, cre], Felix Anker [ctb]

Maintainer: Joerg Polzehl <joerg.polzehl@wias-berlin.de>

References

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- J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.
- J. Polzehl and V. Spokoiny (2006) Propagation-Separation Approach for Local Likelihood Estimation, Prob. Theory and Rel. Fields 135(3), 335-362. DOI:10.1007/s00440-005-0464-1.
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- A. Buades, B. Coll and J. M. Morel (2006). A review of image denoising algorithms, with a new one. Simulation, 4, 490-530. DOI:10.1137/040616024.
- Rudin, L.I., Osher, S. and Fatemi, E. (1992). Nonlinear total variation based noise removal algorithms. Phys. D, 60, 259-268. DOI: 10.1016/0167-2789(92)90242-F.
- Bredies, K., Kunisch, K. and Pock, T. (2010). Total Generalized Variation. SIAM J. Imaging Sci., 3, 492-526. DOI:10.1137/090769521.

Description

Function `gethani` determines a bandwidth that leads to, for the specified kernel, a variance reduction for a non-adaptive kernel estimate by a factor of `value`. `getvofh` calculates the sum of location weights for a given bandwidth vector and kernel. `sofmchi` precomputes the variance of a non-central chi distribution with $2*L$ degrees of freedom as a function of the noncentrality parameter for an interval $c(0, to)$. Functions `residualVariance` and `residualSpatialCorr` are used in package `fmri` to calculate variances and spatial correlations from residual objects.

Usage

```
gethani(x, y, lkern, value, wght, eps = 0.01)
getvofh(bw, lkern, wght)
sofmchi(L, to = 50, delta = 0.01)
residualVariance(residuals, mask, resscale = 1, compact = FALSE)
residualSpatialCorr(residuals, mask, lags = c(5, 5, 3), compact = FALSE)
```

Arguments

<code>x</code>	lower bound of search interval
<code>y</code>	upper bound of search interval
<code>lkern</code>	code for location kernel
<code>value</code>	target sum of location weights
<code>wght</code>	relative size of voxel dimensions $c(0, 0)$ for 1D and $c(w1, 0)$ for 2D problems.
<code>eps</code>	attempted precision for bandwidth search
<code>bw</code>	vector of bandwidths, length equal to 1, 2 or 3 depending on the dimensionality of the problem.
<code>L</code>	number of effective coils, $2*L$ is the degree of freedom of the non-central chi distribution.
<code>to</code>	upper interval bound.
<code>delta</code>	discretization width.
<code>residuals</code>	array of residuals, if <code>compact</code> only containing voxel with mask, otherwise for complete data cubes.
<code>mask</code>	mask of active voxel (e.g. brain masks)
<code>resscale</code>	scale for residuals (residuals may be scaled for optimal integer*2 storage)
<code>compact</code>	logical, determines if only information for voxel within mask or full for full data cubes is given.
<code>lags</code>	positive integer vector of length 3, maximum lags for spatial correlations for each coordinate direction to be computed

Details

These are auxiliary functions not to be used by the user. They are only exported to be available for internal use in packages `fmri`, `dti`, `qMRI` and `adimpro`.

Value

`gethani` returns a vector of bandwidths, `getvofh` returns the variance reduction that would be obtained with a kernel estimate employing the specified kernel and bandwidth, `sofmchi` returns a list with, e.g., components `ncp` and `s2` containing vectors of noncentrality parameter values and corresponding variances, respectively, for the specified noncentral Chi distribution, `residualVariance` returns a vector (`compact==TRUE`) or array (`compact==FALSE`) of voxelwise residual variances, `residualSpatialCorr` returns an array of dimension lags containing spatial correlations.

Note

These functions are for internal use only. They are only exported to be available in other packages.

Author(s)

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aws

AWS for local constant models on a grid

Description

The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient likelihood models on a 1D, 2D or 3D grid. For "Gaussian" models, i.e. regression with additive "Gaussian" errors, a homoskedastic or heteroskedastic model is used depending on the content of `sigma2`

Usage

```
aws(y,hmax=NULL, mask=NULL, aws=TRUE, memory=FALSE, family="Gaussian",
    lkern="Triangle", aggkern="Uniform",
    sigma2=NULL, shape=NULL, scor=0, spmin=0.25,
    ladjust=1,wghts=NULL,u=NULL,graph=FALSE,demo=FALSE,
    testprop=FALSE,maxni=FALSE)
```

Arguments

<code>y</code>	array <code>y</code> containing the observe response (image intensity) data. <code>dim(y)</code> determines the dimensionality and extend of the grid design.
<code>hmax</code>	<code>hmax</code> specifies the maximal bandwidth. Defaults to <code>hmax=250, 12, 5</code> for 1D, 2D, 3D images, respectively. In case of <code>lkern="Gaussian"</code> the bandwidth is assumed to be given in full width half maximum (FWHM) units, i.e., 0.42466 times <code>gridsize</code> .

aws	logical: if TRUE structural adaptation (AWS) is used.
mask	optional logical mask, same dimensionality as y
memory	logical: if TRUE stagewise aggregation is used as an additional adaptation scheme.
family	family specifies the probability distribution. Default is family="Gaussian", also implemented are "Bernoulli", "Poisson", "Exponential", "Volatility", "Variance" and "NCchi". family="Volatility" specifies a Gaussian distribution with expectation 0 and unknown variance. family="Volatility" specifies that $p \cdot y / \theta$ is distributed as χ^2 with $p = \text{shape}$ degrees of freedom. family="NCchi" uses a noncentral Chi distribution with $p = \text{shape}$ degrees of freedom and noncentrality parameter θ
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.
aggkern	character: kernel used in stagewise aggregation, either "Triangle" or "Uniform"
sigma2	sigma2 allows to specify the variance in case of family="Gaussian". Not used if family!="Gaussian". Defaults to NULL. In this case a homoskedastic variance estimate is generated. If $\text{length}(\text{sigma2}) = \text{length}(y)$ then sigma2 is assumed to contain the pointwise variance of y and a heteroscedastic variance model is used.
shape	Allows to specify an additional shape parameter for certain family models. Currently only used for family="Variance", that is χ -Square distributed observations with shape degrees of freedom.
scorr	The vector scorrr allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).
spmin	Determines the form (size of the plateau) in the adaptation kernel. Not to be changed by the user.
ladjust	factor to increase the default value of lambda
wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with $u=0$
graph	If graph=TRUE intermediate results are illustrated after each iteration step. Defaults to graph=FALSE.
demo	If demo=TRUE the function pauses after each iteration. Defaults to demo=FALSE.
testprop	If set this provides diagnostics for testing the propagation condition. The values of y should correspond to the specified family and a global model.
maxni	If TRUE use $\max_{l \leq k} (N_i^{(l)})$ instead of $(N_i^{(k)})$ in the definition of the statistical penalty.

Details

The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient likelihood models on a 1D, 2D or 3D grid. For "Gaussian" models, i.e. regression with additive "Gaussian" errors, a homoskedastic or heteroskedastic model is used depending on the content of `sigma2`. `aws==FALSE` provides the stagewise aggregation procedure from Belomestny and Spokoiny (2004). `memory==FALSE` provides Adaptive weights smoothing without control by stagewise aggregation.

The essential parameter in the procedure is a critical value `lambda`. This parameter has an interpretation as a significance level of a test for equivalence of two local parameter estimates. Optimal values mainly depend on the chosen family. Values set internally are chosen to fulfil a propagation condition, i.e. in case of a constant (global) parameter value and large `hmax` the procedure provides, with a high probability, the global (parametric) estimate. More formally we require the parameter `lambda` to be specified such that $\mathbf{E}|\hat{\theta}^k - \theta| \leq (1 + \alpha)\mathbf{E}|\tilde{\theta}^k - \theta|$ where $\hat{\theta}^k$ is the `aws`-estimate in step `k` and $\tilde{\theta}^k$ is corresponding nonadaptive estimate using the same bandwidth (`lambda=Inf`). The value of `lambda` can be adjusted by specifying the factor `ladjust`. Values `ladjust>1` lead to an less effective adaptation while `ladjust<<1` may lead to random segmentation of, with respect to a constant model, homogeneous regions.

The numerical complexity of the procedure is mainly determined by `hmax`. The number of iterations is approximately $\text{Const} \cdot d \cdot \log(hmax) / \log(1.25)$ with `d` being the dimension of `y` and the constant depending on the kernel `lkern`. Complexity in each iteration step is $\text{Const} \cdot hakt \cdot n$ with `hakt` being the actual bandwidth in the iteration step and `n` the number of design points. `hmax` determines the maximal possible variance reduction.

Value

returns an object of class `aws` with slots

```

y = "numeric"      y
dy = "numeric"     dim(y)
x = "numeric"      numeric(0)
ni = "integer"     integer(0)
mask = "logical"   logical(0)
theta = "numeric"  Estimates of regression function, length: length(y)
mae = "numeric"    Mean absolute error for each iteration step if u was specified, numeric(0) else
var = "numeric"    approx. variance of the estimates of the regression function. Please note that
                    this does not reflect variability due to randomness of weights.
xmin = "numeric"   numeric(0)
xmax = "numeric"   numeric(0)
wghts = "numeric"  numeric(0), ratio of distances wghts[-1]/wghts[1]
degree = "integer" 0

```

```

hmax = "numeric"
          effective hmax
sigma2 = "numeric"
          provided or estimated error variance
scorr = "numeric"
          scor
family = "character"
          family
shape = "numeric"
          shape
lkern = "integer"
          integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
          5="Gaussian"
lambda = "numeric"
          effective value of lambda
ladjust = "numeric"
          effective value of ladjust

aws = "logical"  aws
memory = "logical"
          memory
homogen = "logical"
          homogen
earlystop = "logical"
          FALSE
varmodel = "character"
          "Constant"
vcoef = "numeric"
          numeric(0)
call = "function"
          the arguments of the call to aws

```

Note

use `setCores='number of threads'` to enable parallel execution.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

- J. Polzehl, K. Tabelow (2019). Magnetic Resonance Brain Imaging: Modeling and Data Analysis Using R. Springer, Use R! series. Appendix A. Doi:10.1007/978-3-030-29184-6.
- J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.
- J. Polzehl, V. Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62 , (2000) , pp. 335–354. DOI:10.1111/1467-9868.00235.
- J. Polzehl, V. Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006) , pp. 335–362. DOI:10.1007/s00440-005-0464-1.

See Also

See also [paws](#), [lpaws](#), [vaws](#), [link{awsdata}](#), [aws.irreg](#), [aws.gaussian](#)

Examples

```
require(aws)
# 1D local constant smoothing
## Not run: demo(aws_ex1)
## Not run: demo(aws_ex2)
# 2D local constant smoothing
## Not run: demo(aws_ex3)
```

aws-class

*Class "aws"***Description**

The "aws" class is used for objects obtained by functions `aws`, `lpaws`, `aws.irreg` and `aws.gaussian`.

Objects from the Class

Objects are created by calls to functions `aws`, `lpaws`, `aws.irreg` and `aws.gaussian`.

Slots

.Data: Object of class "list", usually empty.

y: Object of class "array" containing the original (response) data

dy: Object of class "numeric" dimension attribute of y

nvec: Object of class "integer" leading dimension of y in vector valued data.

x: Object of class "numeric" if provided the design points

ni: Object of class "numeric" sum of weights used in final estimate

mask: Object of class "logical" mask of design points where computations are performed

theta: Object of class "array" contains the smoothed object and in case of function `lpaws` its derivatives up to the specified degree. Dimension is $\dim(\theta) = c(dy, p)$

hseq: Sequence of bandwidths employed.

mae: Object of class "numeric" Mean absolute error with respect to array in argument `u` if provided.

psnr: Object of class "numeric" Peak Signal to Noise Ratio (PSNR) with respect to array in argument `u` if provided.

var: Object of class "numeric" pointwise variance of $\theta[\dots, 1]$

xmin: Object of class "numeric" min of x in case of irregular design

xmax: Object of class "numeric" max of x in case of irregular design

wghts: Object of class "numeric" weights used in location penalty for different coordinate directions, corresponds to ratios of distances in coordinate directions 2 and 3 to and distance in coordinate direction 1.

degree: Object of class "integer" degree of local polynomials used in function lpaws

hmax: Object of class "numeric" maximal bandwidth

sigma2: Object of class "numeric" estimated error variance

scorr: Object of class "numeric" estimated spatial correlation

family: Object of class "character" distribution of y, can be any of c("Gaussian", "Bernoulli", "Poisson", "Exponential", "Volatility", "Variance")

shape: Object of class "numeric" possible shape parameter of distribution of y

lkern: Object of class "integer" location kernel, can be any of c("Triangle", "Quadratic", "Cubic", "Plateau", "Gaussian", "Exponential", "Volatility", "Variance") defaults to "Triangle"

lambda: Object of class "numeric" scale parameter used in adaptation

ladjust: Object of class "numeric" factor to adjust scale parameter with respect to its predetermined default.

aws: Object of class "logical" Adaptation by Propagation-Separation

memory: Object of class "logical" Adaptation by Stagewise Aggregation

homogen: Object of class "logical" detect regions of homogeneity (used to speed up the calculations)

earlystop: Object of class "logical" further speedup in function lpaws estimates are fixed if sum of weights does not increase with iterations.

varmodel: Object of class "character" variance model used in function aws.gaussian

vcoef: Object of class "numeric" estimates variance parameters in function aws.gaussian

call: Object of class "call" that created the object.

Methods

extract signature(x = "aws"): ...

risk signature(y = "aws"): ...

plot Method for Function 'plot' in Package 'aws'.

show Method for Function 'show' in Package 'aws'.

print Method for Function 'print' in Package 'aws'.

summary Method for Function 'summary' in Package 'aws'.

Author(s)

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References

Joerg Polzehl, Vladimir Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62, (2000), pp. 335–354

Joerg Polzehl, Vladimir Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006), pp. 335–362.

See Also

[aws](#), [lpaws](#), [aws.irreg](#), [aws.gaussian](#)

Examples

```
showClass("aws")
```

aws.gaussian	<i>Adaptive weights smoothing for Gaussian data with variance depending on the mean.</i>
--------------	--

Description

The function implements an semiparametric adaptive weights smoothing algorithm designed for regression with additive heteroskedastic Gaussian noise. The noise variance is assumed to depend on the value of the regression function. This dependence is modeled by a global parametric (polynomial) model.

Usage

```
aws.gaussian(y, hmax = NULL, hpre = NULL, aws = TRUE, memory = FALSE,
             varmodel = "Constant", lkern = "Triangle",
             aggkern = "Uniform", scor = 0, mask=NULL, ladjust = 1,
             wghts = NULL, u = NULL, varprop = 0.1, graph = FALSE, demo = FALSE)
```

Arguments

y	y contains the observed response data. dim(y) determines the dimensionality and extend of the grid design.
hmax	hmax specifies the maximal bandwidth. Defaults to hmax=250, 12, 5 for dd=1, 2, 3, respectively.
hpre	Describe hpre Bandwidth used for an initial nonadaptive estimate. The first estimate of variance parameters is obtained from residuals with respect to this estimate.
aws	logical: if TRUE structural adaptation (AWS) is used.
memory	logical: if TRUE stagewise aggregation is used as an additional adaptation scheme.
varmodel	Implemented are "Constant", "Linear" and "Quadratic" referring to a polynomial model of degree 0 to 2.
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.

aggkern	character: kernel used in stagewise aggregation, either "Triangle" or "Uniform"
scorr	The vector scorrr allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).
mask	Restrict smoothing to points where mask==TRUE. Defaults to TRUE in all voxel.
ladjust	factor to increase the default value of lambda
wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with u=0
varprop	Small variance estimates are replaced by varprop times the mean variance.
graph	If graph=TRUE intermediate results are illustrated after each iteration step. Defaults to graph=FALSE.
demo	If demo=TRUE the function pauses after each iteration. Defaults to demo=FALSE.

Details

The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient likelihood models on a 1D, 2D or 3D grid. In contrast to function `aws` observations are assumed to follow a Gaussian distribution with variance depending on the mean according to a specified global variance model. `aws==FALSE` provides the stagewise aggregation procedure from Belomestny and Spokoiny (2004). `memory==FALSE` provides Adaptive weights smoothing without control by stagewise aggregation.

The essential parameter in the procedure is a critical value `lambda`. This parameter has an interpretation as a significance level of a test for equivalence of two local parameter estimates. Values set internally are chosen to fulfil a propagation condition, i.e. in case of a constant (global) parameter value and large `hmax` the procedure provides, with a high probability, the global (parametric) estimate. More formally we require the parameter `lambda` to be specified such that $\mathbf{E}|\hat{\theta}^k - \theta| \leq (1 + \alpha)\mathbf{E}|\tilde{\theta}^k - \theta|$ where $\hat{\theta}^k$ is the `aws`-estimate in step `k` and $\tilde{\theta}^k$ is corresponding nonadaptive estimate using the same bandwidth (`lambda=Inf`). The value of `lambda` can be adjusted by specifying the factor `ladjust`. Values `ladjust>1` lead to an less effective adaptation while `ladjust<<1` may lead to random segmentation of, with respect to a constant model, homogeneous regions.

The numerical complexity of the procedure is mainly determined by `hmax`. The number of iterations is approximately $\text{Const} \cdot d \cdot \log(hmax) / \log(1.25)$ with `d` being the dimension of `y` and the constant depending on the kernel `lkern`. Complexity in each iteration step is $\text{Const} \cdot hakt \cdot n$ with `hakt` being the actual bandwidth in the iteration step and `n` the number of design points. `hmax` determines the maximal possible variance reduction.

Value

returns an object of class `aws` with slots

```

y = "numeric"    y
dy = "numeric"   dim(y)
x = "numeric"    numeric(0)
```

```

ni = "integer"    integer(0)
mask = "logical"
                logical(0)
theta = "numeric"
                Estimates of regression function, length: length(y)
mae = "numeric"   Mean absolute error for each iteration step if u was specified, numeric(0) else
var = "numeric"   approx. variance of the estimates of the regression function. Please note that
                this does not reflect variability due to randomness of weights.
xmin = "numeric"
                numeric(0)
xmax = "numeric"
                numeric(0)
wghts = "numeric"
                numeric(0), ratio of distances wghts[-1]/wghts[1]
degree = "integer"
                0
hmax = "numeric"
                effective hmax
sigma2 = "numeric"
                provided or estimated error variance
scorr = "numeric"
                scor
family = "character"
                "Gaussian"
shape = "numeric"
                NULL
lkern = "integer"
                integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
                5="Gaussian"
lambda = "numeric"
                effective value of lambda
ladjust = "numeric"
                effective value of ladjust
aws = "logical"   aws
memory = "logical"
                memory
homogen = "logical"
                homogen
earlystop = "logical"
                FALSE
varmodel = "character"
                varmodel
vcoef = "numeric"
                estimated parameters of the variance model
call = "function"
                the arguments of the call to aws.gaussian

```

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

Joerg Polzehl, Vladimir Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62 , (2000) , pp. 335–354

Joerg Polzehl, Vladimir Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006) , pp. 335–362.

Joerg Polzehl, Vladimir Spokoiny, in V. Chen, C.; Haerdle, W. and Unwin, A. (ed.) Handbook of Data Visualization Structural adaptive smoothing by propagation-separation methods Springer-Verlag, 2008, 471-492

See Also

See also [aws](#), [link{awsdata}](#), [aws.irreg](#)

Examples

```
require(aws)
```

aws.irreg

local constant AWS for irregular (1D/2D) design

Description

The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient Gaussian models on a 1D or 2D irregular design. The function allows for a parametric (polynomial) mean-variance dependence.

Usage

```
aws.irreg(y, x, hmax = NULL, aws=TRUE, memory=FALSE, varmodel = "Constant",
          lkern = "Triangle", aggkern = "Uniform", sigma2 = NULL, nbins = 100,
          hpre = NULL, henv = NULL, ladjust = 1, varprop = 0.1, graph = FALSE)
```

Arguments

y	The observed response vector (length n)
x	Design matrix, dimension n x d, d %in% 1:2
hmax	hmax specifies the maximal bandwidth. Unit is binwidth in the first dimension.
aws	logical: if TRUE structural adaptation (AWS) is used.
memory	logical: if TRUE stagewise aggregation is used as an additional adaptation scheme.

varmodel	determines the model that relates variance to mean. Either "Constant", "Linear" or "Quadratic".
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian"
aggkern	character: kernel used in stagewise aggregation, either "Triangle" or "Uniform"
sigma2	sigma2 allows to specify the variance in case of varmodel="Constant", estimated if not given.
nbins	numer of bins, can be NULL, a positive integer or a vector of positive integers (length d)
hpre	smoothing bandwidth for initial variance estimate
henv	radius of balls around each observed design point where estimates will be calculated
ladjust	factor to increase the default value of lambda
varprop	exclude the largest 100*varprop% squared residuals when estimating the error variance
graph	If graph=TRUE intermediate results are illustrated after each iteration step. Defaults to graph=FALSE.

Details

Data are first binned (1D/2D), then aws is performed on all datapoints within distance $\leq$ henv of nonempty bins.

Value

returns an object of class aws with slots

y = "numeric"	y
dy = "numeric"	dim(y)
x = "numeric"	x
ni = "integer"	number of observations per bin
mask = "logical"	bins where parameters have been estimated
theta = "numeric"	Estimates of regression function, length: length(y)
mae = "numeric"	numeric(0)
var = "numeric"	approx. variance of the estimates of the regression function. Please note that this does not reflect variability due to randomness of weights.
xmin = "numeric"	vector of minimal x-values (bins)
xmax = "numeric"	vector of maximal x-values (bins)
wghts = "numeric"	relative binwidths

```

degree = "integer"
      0
hmax = "numeric"
      effective hmax
sigma2 = "numeric"
      provided or estimated error variance
scorr = "numeric"
      0
family = "character"
      "Gaussian"
shape = "numeric"
      numeric(0)
lkern = "integer"
      integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
      5="Gaussian"
lambda = "numeric"
      effective value of lambda
ladjust = "numeric"
      effective value of ladjust

aws = "logical"  aws
memory = "logical"
      memory
homogen = "logical"
      FALSE
earlystop = "logical"
      FALSE
varmodel = "character"
      varmodel
vcoef = "numeric"
      estimated coefficients in variance model
call = "function"
      the arguments of the call to aws

```

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>

References

J. Polzehl, V. Spokoiny, in V. Chen, C.; Haerdle, W. and Unwin, A. (ed.) Handbook of Data Visualization Structural adaptive smoothing by propagation-separation methods. Springer-Verlag, 2008, 471-492. DOI:10.1007/978-3-540-33037-0_19.

See Also

See also [lpaws](#), [link{awsdata}](#), [lpaws](#)

Examples

```
require(aws)
# 1D local constant smoothing
## Not run: demo(irreg_ex1)
# 2D local constant smoothing
## Not run: demo(irreg_ex2)
```

aws.segment

Segmentation by adaptive weights for Gaussian models.

Description

The function implements a modification of the adaptive weights smoothing algorithm for segmentation into three classes. The

Usage

```
aws.segment(y, level, delta = 0, hmax = NULL, hpre = NULL, mask = NULL,
            varmodel = "Constant", lkern = "Triangle", scorr = 0, ladjust = 1,
            wghts = NULL, u = NULL, varprop = 0.1, ext = 0, graph = FALSE,
            demo = FALSE, fov=NULL)
```

Arguments

y	y contains the observed response data. <code>dim(y)</code> determines the dimensionality and extend of the grid design.
level	center of second class
delta	half width of second class
hmax	hmax specifies the maximal bandwidth. Defaults to <code>hmax=250, 12, 5</code> for <code>dd=1, 2, 3</code> , respectively.
hpre	Describe hpre Bandwidth used for an initial nonadaptive estimate. The first estimate of variance parameters is obtained from residuals with respect to this estimate.
mask	optional logical mask, same dimensionality as y
varmodel	Implemented are "Constant", "Linear" and "Quadratic" referring to a polynomial model of degree 0 to 2.
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.
scorr	The vector <code>scorr</code> allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).

ladjust	factor to increase the default value of lambda
wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with $u=0$
varprop	Small variance estimates are replaced by varprop times the mean variance.
ext	Intermediate results are fixed if the test statistics exceeds the critical value by ext.
graph	If graph=TRUE intermediate results are illustrated after each iteration step. Defaults to graph=FALSE.
demo	If demo=TRUE the function pauses after each iteration. Defaults to demo=FALSE.
fov	Field of view. Size of region (sample size) to adjust for in multiscale testing.

Details

The image is segmented into three parts by performing multiscale tests of the hypotheses $H1$ value $\geq \text{level} - \delta$ and $H2$ value $\leq \text{level} + \delta$. Pixel where the first hypothesis is rejected are classified as -1 (segment 1) while rejection of $H2$ results in classification 1 (segment 3). Pixel where neither $H1$ or $H2$ are rejected are assigned to a value 0 (segment 2). Critical values for the tests are adjusted for smoothness at the different scales inspected in the iteration process using results from multiscale testing, see e.g. Duembgen and Spokoiny (2001). Critical values also depend on the size of the region of interest specified in parameter fov.

Within segment 2 structural adaptive smoothing is performed while if a pair of pixel belongs to segment 1 or segment 3 the corresponding weight will be nonadaptive.

Value

returns an object of class aws with slots

y = "numeric"	y
dy = "numeric"	dim(y)
x = "numeric"	numeric(0)
ni = "integer"	integer(0)
mask = "logical"	logical(0)
segment = "integer"	Segmentation results, class numbers 1-3
theta = "numeric"	Estimates of regression function, length: length(y)
mae = "numeric"	Mean absolute error for each iteration step if u was specified, numeric(0) else
var = "numeric"	approx. variance of the estimates of the regression function. Please note that this does not reflect variability due to randomness of weights.
xmin = "numeric"	numeric(0)

```

xmax = "numeric"
        numeric(0)
wghts = "numeric"
        numeric(0), ratio of distances wghts[-1]/wghts[1]
degree = "integer"
        0
hmax = "numeric"
        effective hmax
sigma2 = "numeric"
        provided or estimated error variance
scorr = "numeric"
        scor
family = "character"
        "Gaussian"
shape = "numeric"
        NULL
lkern = "integer"
        integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
        5="Gaussian"
lambda = "numeric"
        effective value of lambda
ladjust = "numeric"
        effective value of ladjust

aws = "logical"  aws
memory = "logical"
        memory
homogen = "logical"
        FALSE
earlystop = "logical"
        FALSE
varmodel = "character"
        varmodel
vcoef = "numeric"
        estimated parameters of the variance model
call = "function"
        the arguments of the call to aws.gaussian

```

Note

This function is still experimental and may be changes considerably in future.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

J. Polzehl, H.U. Voss, K. Tabelow (2010). Structural adaptive segmentation for statistical parametric mapping, *NeuroImage*, 52, pp. 515–523. DOI:10.1016/j.neuroimage.2010.04.241

Duembgen, L. and Spokoiny, V. (2001). Multiscale testing of qualitative hypotheses. *Ann. Stat.* 29, 124–152.

Polzehl, J. and Spokoiny, V. (2006). Propagation-Separation Approach for Local Likelihood Estimation. *Probability Theory and Related Fields.* 3 (135) 335 - 362. DOI:10.1007/s00440-005-0464-1

See Also

[aws](#), [aws.gaussian](#)

Examples

```
require(aws)
```

awsdata	<i>Extract information from an object of class aws</i>
---------	--

Description

Extract data and estimates from an object of class aws

Usage

```
awsdata(awsobj, what)
```

Arguments

awsobj	an object of class aws
what	can be "data" (extracts observed response), "theta" (estimated parameters), "est" (estimated regression function), "var" (approx. variance of estimated regression function), "sd" (approx. standard deviation of estimated regression function), "sigma2" (error variance), "mae" (mean absolute error for each iteration step, if available), "ni" (number of observations per bin), "mask" (logical indicator for bins where the regression function is estimated). "bi" (array of sum of weights or NULL) "bi2" (array of sum of squared weights or NULL)

Details

The returned object is formatted as an array if appropriate. The returned object may be NULL if the information is not available.

Value

an vector or array containing the specified information.

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

Joerg Polzehl, Vladimir Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62 , (2000) , pp. 335–354

Joerg Polzehl, Vladimir Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006) , pp. 335–362.

Joerg Polzehl, Vladimir Spokoiny, in V. Chen, C.; Haerdle, W. and Unwin, A. (ed.) Handbook of Data Visualization Structural adaptive smoothing by propagation-separation methods Springer-Verlag, 2008, 471-492

See Also

`link{awsdata}`, [aws](#), [aws.irreg](#)

Examples

```
require(aws)
# 1D local constant smoothing
## Not run: demo(aws_ex1)
## Not run: demo(aws_ex2)
# 2D local constant smoothing
## Not run: demo(aws_ex3)
# 1D local polynomial smoothing
## Not run: demo(lpaws_ex1)
# 2D local polynomial smoothing
## Not run: demo(lpaws_ex2)
# 1D irregular design
## Not run: demo(irreg_ex1)
# 2D irregular design
## Not run: demo(irreg_ex2)
```

awsLocalSigma

3D variance estimation

Description

Functions for 3D variance estimation. `awsLocalSigma` implements the local adaptive variance estimation procedure introduced in Tabelow, Voss and Polzehl (2015). `awsLinsd` uses a parametric model for variance/mean dependence. Functions `AFLocalSigma` and `estGlobalSigma` implement various proposals for local and global variance estimates from Aja-Fernandez (2009, 2013) and a global variant of the approach from Tabelow, Voss and Polzehl (2015).

Usage

```

awsLocalSigma(y, steps, mask, ncoils, vext = c(1, 1), lambda = 5,
  minni = 2, hsig = 5, sigma = NULL, family = c("NCchi", "Gauss"),
  verbose = FALSE, trace = FALSE, u = NULL)
awslinsd(y, hmax = NULL, hpre = NULL, h0 = NULL, mask = NULL,
  ladjust = 1, wghts = NULL, varprop = 0.1, A0, A1)
AFLocalSigma(y, ncoils, level = NULL, mask = NULL, h = 2, hadj = 1,
  vext = c(1, 1))
estGlobalSigma(y, mask = NULL, ncoils = 1, steps = 16, vext = c(1, 1),
  lambda = 20, hinit = 2, hadj = 1, q = 0.25, level = NULL,
  sequence = FALSE, method = c("awsVar", "awsMAD", "AFmodevn",
    "AFmodem1chi", "AFbkm2chi", "AFbkm1chi"))
estimateSigmaCompl(magnitude, phase, mask, kstar = 20, kmin = 8, hsig = 5,
  lambda = 12, verbose = TRUE)

```

Arguments

y	3D array of image intensities.
steps	number of steps in adaptive weights smoothing, used to reveal the underlying mean structure.
mask	restrict computations to voxel in mask, if <code>is.null(mask)</code> all voxel are used. In function <code>estGlobalSigma</code> mask should refer to background for method <code>%in% c("modem1chi", "bkm2chi", "bkm1chi")</code> and to voxel within the head for method <code>=="modevn"</code> .
ncoils	effective number of coils, or equivalently number of effective degrees of freedom of non-central chi distribution divided by 2.
vext	voxel extensions or relative voxel extensions
lambda	scale parameter in adaptive weights smoothing
minni	minimal bandwidth for calculating local variance estimates
hsig	bandwidth for median filter
sigma	optional initial global variance estimate
family	type of distribution, either noncentral Chi ("NCchi") or Gaussian ("Gauss")
verbose	if <code>verbose==TRUE</code> density plots and quantiles of local estimates of sigma are provided.
trace	if <code>trace==TRUE</code> intermediate results for each step are returned in component <code>ters</code> for all voxel in mask.
u	if <code>verbose==TRUE</code> an array of noncentrality parameters for comparisons. Internal use for tests only
hmax	maximal bandwidth
hpre	minimal bandwidth
h0	bandwidth vector characterizing to spatial correlation as correlation induced by convolution with a Gaussian kernel
ladjust	correction factor for lambda
wghts	relative voxel extensions

varprop	defines a lower bound for the estimated variance as $\text{varprop} \times \text{mean}(\sigma^2_{\text{hat}}$
A0	select voxel with $A_0 < \theta < A_1$ to estimate parameters of the variance model
A1	select voxel with $A_0 < \theta < A_1$ to estimate parameters of the variance model
level	threshold for mask definition
h	bandwidth for local variance estimates.
hinit	minimal bandwidth for local variance estimates with method="awsxxx".
hadj	bandwidth for mode estimation
q	Quantile for interquantile estimate of standard deviation
sequence	logical, return sequence of estimated variances for iterative methods.
method	determines variance estimation method
magnitude	magnitude of complex 3D image
phase	phase of complex 3D image
kstar	number of steps in adaptive weights smoothing, used to reveal the underlying mean structure.
kmin	iteration to start adaptation

Value

all functions return lists with variance estimates in component sigma

Author(s)

Jörg Polzehl <polzehl@wias-berlin.de>

References

- K. Tabelow, H.U. Voss, J. Polzehl, Local estimation of the noise level in MRI using structural adaptation, *Medical Image Analysis*, 20 (2015), 76–86. DOI:10.1016/j.media.2014.10.008.
- S. Aja-Fernandez, V. Brion, A. Tristan-Vega, Effective noise estimation and filtering from correlated multiple-coil MR data. *Magn Reson Imaging*, 31 (2013), 272–285. DOI:10.1016/j.mri.2012.07.006
- S. Aja-Fernandez, A. Tristan-Vega, C. Alberola-Lopez, Noise estimation in single- and multiple-coil magnetic resonance data based on statistical models. *Magn Reson Imaging*, 27 (2009), 1397–1409. DOI:10.1016/j.mri.2009.05.025.

awssegment-class	Class "awssegment"
------------------	--------------------

Description

The "aws" class is used for objects obtained by functions `aws.segment`

Objects from the Class

Objects are created by calls to functions `aws.segment`

Slots

.Data: Object of class "list", usually empty.

y: Object of class "array" containing the original (response) data

dy: Object of class "numeric" dimension attribute of y

x: Object of class "numeric" if provided the design points

ni: Object of class "numeric" sum of weights used in final estimate

mask: Object of class "logical" mask of design points where computations are performed

segment: Object of class "array" segmentation results (3 segments coded by $c(-1, 0, 1)$)

level: Object of class "numeric" center of segment 0

delta: Object of class "numeric" half width of segment 0

theta: Object of class "array" ~~

theta: Object of class "array" contains the smoothed object and in case of function `lpaws` its derivatives up to the specified degree. Dimension is $\dim(\text{theta})=c(dy, p)$

mae: Object of class "numeric" Mean absolute error with respect to array in argument `u` if provided.

var: Object of class "numeric" pointwise variance of `theta[... ,1]`

xmin: Object of class "numeric" not used

xmax: Object of class "numeric" not used

wghts: Object of class "numeric" weights used in location penalty for different coordinate directions

degree: not used

hmax: Object of class "numeric" maximal bandwidth

sigma2: Object of class "numeric" estimated error variance

scorr: Object of class "numeric" estimated spatial correlation

family: Object of class "character" distribution of y, can be any of `c("Gaussian", "Bernoulli", "Poisson", "Exponential", "Volatility", "Variance")`

shape: Object of class "numeric" possible shape parameter of distribution of y

lkern: Object of class "integer" location kernel, can be any of c("Triangle", "Quadratic", "Cubic", "Plateau", "Gaussian") defaults to "Triangle"

lambda: Object of class "numeric" scale parameter used in adaptation

ladjust: Object of class "numeric" factor to adjust scale parameter with respect to its predetermined default.

aws: Object of class "logical" Adaptation by Propagation-Separation

memory: Object of class "logical" Adaptation by Stagewise Aggregation

homogen: Object of class "logical" detect regions of homogeneity (used to speed up the calculations) currently FALSE

earlystop: Object of class "logical" currently FALSE

varmodel: Object of class "character" variance model used currently "Gaussian"

vcoef: Object of class "numeric" contains NULL

call: Object of class "call" that created the object.

Methods

extract signature(x = "awssegment"): ...

plot signature(x = "awssegment"): ...

print signature(x = "awssegment"): ...

risk signature(y = "awssegment"): ...

show signature(object = "awssegment"): ...

summary signature(object = "awssegment"): ...

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>

See Also

[aws.segment](#)

Examples

```
showClass("awssegment")
```

awstestprop

*Propagation condition for adaptive weights smoothing***Description**

The function enables testing of the propagation condition in order to select appropriate values for the parameter `lambda` in function `aws`.

Usage

```
awstestprop(dy, hmax, theta = 1, family = "Gaussian", lkern = "Triangle",
            aws = TRUE, memory = FALSE, shape = 2, homogeneous=TRUE, varadapt=FALSE,
            ladjust = 1, spmin=0.25, seed = 1, minlevel=1e-6, maxz=25, diffz=.5,
            maxni=FALSE, verbose=FALSE)
pawstestprop(dy, hmax, theta = 1, family = "Gaussian", lkern = "Triangle",
             aws = TRUE, patchsize=1, shape = 2,
             ladjust = 1, spmin = 0.25, seed = 1, minlevel = 1e-6,
             maxz = 25, diffz = .5, maxni = FALSE, verbose = FALSE)
```

Arguments

<code>dy</code>	Dimension of grid used in 1D, 2D or 3D. May also be specified as an array of values. In this case data are generated with parameters $\text{dy} - \text{mean}(\text{dy}) + \text{theta}$ and the propagation condition is tested as if <code>theta</code> is the true parameter. This can be used to study properties for a slightly misspecified structural assumption.
<code>hmax</code>	Maximum bandwidth.
<code>theta</code>	Parameter determining the distribution in case of family <code>c("Poisson", "Bernoulli")</code>
<code>family</code>	family specifies the probability distribution. Default is <code>family="Gaussian"</code> , also implemented are "Bernoulli", "Poisson", "Exponential", "Volatility", "Variance" and "NCchi". <code>family="Volatility"</code> specifies a Gaussian distribution with expectation 0 and unknown variance. <code>family="Volatility"</code> specifies that $p*y/\text{theta}$ is distributed as χ^2 with $p=\text{shape}$ degrees of freedom. <code>family="NCchi"</code> uses a noncentral Chi distribution with $p=\text{shape}$ degrees of freedom and noncentrality parameter <code>theta</code> .
<code>lkern</code>	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian"
<code>aws</code>	logical: if TRUE structural adaptation (AWS) is used.
<code>patchsize</code>	patchsize in case of paws.
<code>memory</code>	logical: if TRUE stagewise aggregation is used as an additional adaptation scheme.
<code>shape</code>	Allows to specify an additional shape parameter for certain family models. Currently only used for <code>family="Variance"</code> , that is χ -Square distributed observations with <code>shape</code> degrees of freedom.

homogeneous	if homogeneous==FALSE and family==Gaussian then create heterogeneous variances according to a chi-squared distribution with number of degrees of freedom given by sphere
varadapt	if varadapt==TRUE use inverse of variance reduction instead of sum of weights in definition of statistical penalty.
ladjust	Factor to increase the default value of lambda
spmin	Determines the form (size of the plateau) in the adaptation kernel. Not to be changed by the user.
seed	Seed value for random generator.
minlevel	Minimum exceedence probability to use in contour plots.
maxz	Maximum of z-scale in plots.
diffz	Gridlength in z
maxni	If TRUE use $\max_{l \leq k} (N_i^{(l)})$ instead of $(N_i^{(k)})$ in the definition of the statistical penalty.
verbose	If TRUE provide additional information.

Details

Estimates exceedence probabilities

Results for intermediate steps are provided as contour plots. For a good choice of lambda (ladjust) the contours up to probabilities of 1e-5 should be vertical.

Value

A list with components

h	Sequence of bandwidths used
z	seq(0, 30, .5), the quantiles exceedence probabilities refer to
prob	the matrix of exceedence probabilities, columns corresponding to h
probna	the matrix of exceedence probabilities for corresponding nonadaptive estimates, columns corresponding to h

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

S. Becker, P. Mathe, Electron. J. Statist. (2013), 2702-2736, doi:10.1214/13-EJS860

See Also

[aws](#)

awsweights

*Generate weight scheme that would be used in an additional aws step***Description**

Utility function to create a weighting scheme for an additional aws step. Inteded to be used for illustrations only.

Usage

```
awsweights(awsobj, spmin = 0.25, inx = NULL)
```

Arguments

awsobj	object obtained by a call to function aws
spmin	Size of the plateau in the adaptation kernel.
inx	either a matrix of dimension $\text{length}(\text{awsobj@dy}) \times \text{number of points containing the integer coordinates of points of interest}$ or NULL. In the latter case the weight scheme for all points is generated.

Value

an array of either dimension $\text{awsobj@dy} \times \text{number of points}$ or $\text{awsobj@dy} \times \text{awsobj@dy}$

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

Joerg Polzehl, Vladimir Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62 , (2000) , pp. 335–354

Joerg Polzehl, Vladimir Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006) , pp. 335–362.

Joerg Polzehl, Kostas Papafitsoros, Karsten Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.

See Also

See also [aws](#)

binning

Binning in 1D, 2D or 3D

Description

The function performs a binning in 1D, 2D or 3D.

Usage

```
binning(x, y, nbins, xrange = NULL)
```

Arguments

x	design matrix, dimension $n \times d$, $d \%in\% 1:3$.
y	either a response vector of length n or NULL
nbins	vector of length d containing number of bins for each dimension, may be set to NULL
xrange	range for endpoints of bins for each dimension, either matrix of dimension $2 \times d$ or NULL. xrange is increased if the cube defined does not contain all design points.

Value

A list with components

x	matrix of coordinates of non-empty bin centers
x.freq	number of observations in nonempty bins
midpoints.x1	Bin centers in dimension 1
midpoints.x2	if $d > 1$ Bin centers in dimension 2
midpoints.x3	if $d > 2$ Bin centers in dimension 3
breaks.x1	Break points dimension 1
breaks.x2	if $d > 1$ Break points dimension 2
breaks.x3	if $d > 2$ Break points dimension 3
table.freq	number of observations per bin
means	if <code>!is.null(y)</code> mean of y in non-empty bins
devs	if <code>!is.null(y)</code> standard deviations of y in non-empty bins

Note

This function has been adapted from the code of function binning in package sm.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>

See Also

See Also as [aws.irreg](#)

 extract-methods

Methods for Function extract in Package aws

Description

The method extract and/or compute specified statistics from object of class "aws", "awssegment", ICISmooth and "kernsm".

Usage

```
## S4 method for signature 'aws'
extract(x, what="y")
## S4 method for signature 'awssegment'
extract(x, what="y")
## S4 method for signature 'ICISmooth'
extract(x, what="y")
## S4 method for signature 'kernsm'
extract(x, what="y")
```

Arguments

x	object
what	Statistics to extract, defaults to what="y" corresponding to the original data (response variable). Alternatives are what="yhat" for the smoothed response, what="vhat" for the estimated variance of the smoothed response, what="sigma2" for the estimated error variance of the original data, what="vred" for the variance reduction achieved and in case of signature(x = "ICISmooth") what="hbest" for the selected bandwidth. A vector of any of these choices may be provided.

Methods

signature(x = "ANY") Returns a message that method extract is not defined.

signature(x = "aws") Returns a list with components containing the requested statistics. Component names correspond to tolower(what)

signature(x = "awssegment") Returns a list with components containing the requested statistics. Component names correspond to tolower(what)

signature(x = "ICISmooth") Returns a list with components containing the requested statistics. Component names correspond to tolower(what).

signature(x = "kernsm") Returns a list with components containing the requested statistics. Component names correspond to tolower(what).

ICIcombined	<i>Adaptive smoothing by Intersection of Confidence Intervals (ICI) using multiple windows</i>
-------------	--

Description

The function performs adaptive smoothing by Intersection of Confidence Intervals (ICI) using multiple windows as described in Katkovnik et al (2006)

Usage

```
ICIcombined(y, hmax, hinc = 1.45, thresh = NULL, kern = "Gaussian", m = 0,
            sigma = NULL, nsector = 1, symmetric = FALSE, presmooth = FALSE,
            combine = "weighted", unit = c("SD", "FWHM"))
```

Arguments

y	Object of class "array" containing the original (response) data on a grid
hmax	maximum bandwidth
hinc	factor used to increase the bandwidth from scale to scale
thresh	threshold used in tests to determine the best scale
kern	Determines the kernel function. Object of class "character" kernel, can be any of c("Gaussian", "Uniform", "Triangle", "Epanechnikov", "Biweight", "Triweight"). Defaults to kern="Gaussian".
m	Object of class "integer" vector of length length(dy) determining the order of derivatives specified for the coordinate directions.
sigma	error standard deviation
nsector	number of sectors to use.
symmetric	Object of class "logical" determines if sectors are symmetric with respect to the origin.
presmooth	Object of class "logical" determines if bandwidths are smoothed for more stable results.
combine	Either "weighted" or "minvar". Determines how whether to combine sectorial results a weighted (with inverse variance) mean or to chose the sectorial estimate with minimal variance.
unit	How should the bandwidth be interpreted in case of a Gaussian kernel. For "SD" the bandwidth refers to the standard deviation of the kernel while "FWHM" interprets the bandwidth in terms of Full Width Half Maximum of the kernel.

Details

This mainly follows Chapter 6.2 in Katkovnik et al (2006).

Value

An object of class ICIsmooth

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.

V. Katkovnik, K. Egiazarian and J. Astola, *Local Approximation Techniques in Signal And Image Processing*, SPIE Society of Photo-Optical Instrumentation Engin., 2006, PM157

See Also

[ICIsmooth](#), [ICIsmooth-class](#), [kernsm](#)

ICIsmooth	<i>Adaptive smoothing by Intersection of Confidence Intervals (ICI)</i>
-----------	---

Description

The function performs adaptive smoothing by Intersection of Confidence Intervals (ICI) as described in Katkovnik et al (2006)

Usage

```
ICIsmooth(y, hmax, hinc = 1.45, thresh = NULL, kern = "Gaussian", m = 0,
          sigma = NULL, nsector = 1, sector = 1, symmetric = FALSE,
          presmooth = FALSE, unit = c("SD", "FWHM"))
```

Arguments

y	Object of class "array" containing the original (response) data on a grid
hmax	maximum bandwidth
hinc	factor used to increase the bandwidth from scale to scale
thresh	threshold used in tests to determine the best scale
kern	Determines the kernel function. Object of class "character" kernel, can be any of c("Gaussian", "Uniform", "Triangle", "Epanechnikov", "Biweight", "Triweight"). Defaults to kern="Gaussian".
m	Object of class "integer" vector of length length(dy) determining the order of derivatives specified for the coordinate directions.
sigma	error standard deviation

nsector	number of sectors to use. Positive weights are restricted to the sector selected by sector
sector	Object of class "integer" between 1 and nsector. sector used.
symmetric	Object of class "logical" determines if sectors are symmetric with respect to the origin.
presmooth	Object of class "logical" determines if bandwidths are smoothed for more stable results.
unit	How should the bandwidth be interpreted in case of a Gaussian kernel. For "SD" the bandwidth refers to the standard deviation of the kernel while "FWHM" interprets the bandwidth in terms of Full Width Half Maximum of the kernel.

Details

This mainly follows Chapter 6.1 in Katkovnik et al (2006).

Value

An object of class ICISmooth

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.

V. Katkovnik, K. Egiazarian and J. Astola, *Local Approximation Techniques in Signal And Image Processing*, SPIE Society of Photo-Optical Instrumentation Engin., 2006, PM157

See Also

[ICIcombined](#), [ICISmooth-class](#), [kernsm](#)

ICISmooth-class	Class "ICISmooth"
-----------------	-------------------

Description

The "ICISmooth" class is used for objects obtained by functions ICISmooth and ICIcombined.

Objects from the Class

Objects can be created by calls of the form new("ICISmooth", ...) or by functions ICISmooth and ICIcombined.

Slots

- .Data:** Object of class "list", usually empty.
- y:** Object of class "array" containing the original (response) data
- dy:** Object of class "numeric" dimension attribute of y
- x:** Object of class "numeric" if provided the design points
- hmax:** Object of class "numeric" maximum bandwidth
- hinc:** Object of class "numeric" initial bandwidth
- thresh:** Object of class "numeric" threshold used for bandwidth selection
- kern:** Object of class "character" kernel, can be any of c("Gaussian", "Uniform", "Triangle", "Epanechnikov", "Bivariate")
Defaults to kern="Gaussian".
- m:** Object of class "integer" vector of length length(dy) determining the order of derivatives
specified for the coordinate directios.
- nsector:** Object of class "integer" number of sectors to use.
- sector:** Object of class "integer" sector used.
- symmetric:** Object of class "logical" sectors are symmetric with respect to the origin.
- yhat:** Object of class "array" smoothed response variable
- vhat:** Object of class "array" estimated variance of smoothed response variable
- hbest:** Object of class "array" selected bandwidth(s)
- sigma:** Object of class "numeric" estimated standard deviation of errors in y
- call:** Object of class "call" that created the object.

Methods

extract signature(x = "ICIsMOOTH"): ...
risk signature(y = "ICIsMOOTH"): ...
plot Method for Function 'plot' in Package 'aws'.
show Method for Function 'show' in Package 'aws'.
print Method for Function 'print' in Package 'aws'.
summary Method for Function 'summary' in Package 'aws'.

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

V. Katkovnik, K. Egiazarian and J. Astola, *Local Approximation Techniques in Signal And Image Processing*, SPIE Society of Photo-Optical Instrumentation Engn., 2006, PM157

See Also

ICIs, ICIs, kernsm, aws

Examples

```
showClass("ICISmooth")
```

kernsm

Kernel smoothing on a 1D, 2D or 3D grid

Description

Performs Kernel smoothing on a 1D, 2D or 3D grid by fft

Usage

```
kernsm(y, h = 1, kern = "Gaussian", m = 0, nsector = 1, sector = 1,
       symmetric = FALSE, unit = c("SD", "FWHM"))
```

Arguments

y	Object of class "array" containing the original (response) data on a grid
h	bandwidth
kern	Determines the kernel function. Object of class "character" kernel, can be any of c("Gaussian", "Uniform", "Triangle", "Epanechnikov", "Biweight", "Triweight"). Defaults to kern="Gaussian"
m	Object of class "integer" vector of length length(dy) determining the order of derivatives specified for the coordinate directions.
nsector	number of sectors to use. Positive weights are restricted to the sector selected by sector
sector	Object of class "integer" between 1 and nsector. sector used.
symmetric	Object of class "logical" determines if sectors are symmetric with respect to the origin.
unit	How should the bandwidth be interpreted in case of a Gaussian kernel. For "SD" the bandwidth refers to the standard deviation of the kernel while "FWHM" interprets the bandwidth in terms of Full Width Half Maximum of the kernel.

Details

In case of any ($m > 0$) derivative kernels are generated and applied for the corresponding coordinate directions. If $nsector > 1$ the support of the kernel is restricted to a circular sector determined by sector.

Value

An object of class kernsm

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06 .

V. Katkovnik, K. Egiazarian and J. Astola, *Local Approximation Techniques in Signal And Image Processing*, SPIE Society of Photo-Optical Instrumentation Engin., 2006, PM157

See Also

[kernsm-class](#), [ICIsMOOTH](#), [ICICOMBINED](#)

kernsm-class	Class "kernsm"
--------------	----------------

Description

This class refers to objects created by function kernsm. These objects contain

Objects from the Class

Objects can be created by calls of the form `new("kernsm", ...)`. they are usually created by a call to `function{kernsm}`.

Slots

`.Data`: Object of class "list", usually empty.

`y`: Object of class "array" containing the response in nonparametric regression. The design is assumed to be a 1D, 2D or 3D grid, with dimensionality determined by `dim(y)`.

`dy`: Object of class "numeric" containing `dim(y)`.

`x`: Object of class "numeric" currently not used.

`h`: Object of class "numeric" containing the bandwidth employed.

`kern`: Object of class "character" determining the kernel that was used, can be one of `c("Gaussian", "Uniform", "Triang`

`m`: Object of class "integer" with length `length(dy)` determining the order of derivatives in the corresponding coordinate directions. If `m[i6>0]` a derivative kernel derived from `kern` has been used for the corresponding coordinate direction.

`nsector`: Object of class "integer". If `nsector>1` positive weights are restricted to a segment of a circle (1D or 2D only). The segment is given by `sector`.

`sector`: Object of class "integer" containing the number of the segment used in case of `nsector>1`

`symmetric`: Object of class "logical" determines if the sector is mirrored at the origin.

`yhat`: Object of class "array" with same size and dimension as `y` providing the convolution of `y` with the chosen kernel.

`vred`: Object of class "array" Variance reduction achieved by convolution assuming independence.

`call`: Object of class "function", call that created the object.

Methods**extract** signature(x = "aws"): ...**risk** signature(y = "aws"): ...**plot** Method for Function 'plot' in Package 'aws'.**show** Method for Function 'show' in Package 'aws'.**print** Method for Function 'print' in Package 'aws'.**summary** Method for Function 'summary' in Package 'aws'.**Author(s)**

Jl"org Polzehl <polzehl@wias-berlin.de>

See Also[kernsm](#), [ICIsMOOTH](#), [ICIscombined](#), [ICIsMOOTH](#)**Examples**

showClass("kernsm")

lpaws*Local polynomial smoothing by AWS*

Description

The function allows for structural adaptive smoothing using a local polynomial (degree ≤ 2) structural assumption. Response variables are assumed to be observed on a 1 or 2 dimensional regular grid.

Usage

```
lpaws(y, degree = 1, hmax = NULL, aws = TRUE, memory = FALSE, lkern = "Triangle",
      homogen = TRUE, earlystop = TRUE, aggkern = "Uniform", sigma2 = NULL,
      hw = NULL, ladjust = 1, u = NULL, graph = FALSE, demo = FALSE)
```

Arguments

y	Response, either a vector (1D) or matrix (2D). The corresponding design is assumed to be a regular grid in 1D or 2D, respectively.
degree	Polynomial degree of the local model
hmax	maximal bandwidth
aws	logical: if TRUE structural adaptation (AWS) is used.
memory	logical: if TRUE stagewise aggregation is used as an additional adaptation scheme.

lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.
homogen	logical: if TRUE the function tries to determine regions where weights can be fixed to 1. This may increase speed.
earlystop	logical: if TRUE the function tries to determine points where the homogeneous region is unlikely to change in further steps. This may increase speed.
aggkern	character: kernel used in stagewise aggregation, either "Triangle" or "Uniform"
sigma2	Error variance, the value is estimated if not provided.
hw	Regularisation bandwidth, used to prevent from unidentifiability of local estimates for small bandwidths.
ladjust	factor to increase the default value of lambda
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with $u=0$
graph	logical: If TRUE intermediate results are illustrated graphically. May significantly slow down the computations in 2D. Please avoid using the default X11() on systems build with cairo, use X11(type="Xlib") instead (faster by a factor of 30).
demo	logical: if TRUE wait after each iteration

Value

returns an object of class `aws` with slots

```

y = "numeric"    y
dy = "numeric"   dim(y)
x = "numeric"    numeric(0)
ni = "integer"   integer(0)
mask = "logical" logical(0)
theta = "numeric"
Estimates of regression function and derivatives, length: length(y)*(degree+1)
mae = "numeric"  Mean absolute error for each iteration step if u was specified, numeric(0) else
var = "numeric"  approx. variance of the estimates of the regression function. Please note that
this does not reflect variability due to randomness of weights.
xmin = "numeric" numeric(0)
xmax = "numeric" numeric(0)
wghts = "numeric" numeric(0), ratio of distances wghts[-1]/wghts[1]

```

```

degree = "integer"
    degree
hmax = "numeric"
    effective hmax
sigma2 = "numeric"
    provided or estimated error variance
scorr = "numeric"
    0
family = "character"
    "Gaussian"
shape = "numeric"
    numeric(0)
lkern = "integer"
    integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
    5="Gaussian"
lambda = "numeric"
    effective value of lambda
ladjust = "numeric"
    effective value of ladjust
aws = "logical"  aws
memory = "logical"
    memory
homogen = "logical"
    homogen
earlystop = "logical"
    eralustop
varmodel = "character"
    "Constant"
vcoef = "numeric"
    numeric(0)
call = "function"
    the arguments of the call to lpaws

```

Note

If you specify `graph=TRUE` for 2D problems avoid using the default `X11()` on systems build with `cairo`, use `X11(type="Xlib")` instead (faster by a factor of 30).

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

- J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06 .
- J. Polzehl, V. Spokoiny, in V. Chen, C.; Haerdle, W. and Unwin, A. (ed.) Handbook of Data Visualization Structural adaptive smoothing by propagation-separation methods. Springer-Verlag, 2008, 471-492. DOI:10.1007/978-3-540-33037-0_19.

See Also

`link{awsdata},`[aws](#), [aws.irreg](#)

Examples

```
library(aws)
# 1D local polynomial smoothing
## Not run: demo(lpaws_ex1)
# 2D local polynomial smoothing
## Not run: demo(lpaws_ex2)
```

nlmeans

NLMeans filter in 1D/2D/3D

Description

Implements the Non-Local-Means Filter of Buades et al 2005

Usage

```
nlmeans(x, lambda, sigma, patchhw = 1, searchhw = 7, pd = NULL)
```

Arguments

<code>x</code>	1, 2 or 3-dimensional array of observed response (image intensity) data.
<code>lambda</code>	scale factor for kernel in image space.
<code>sigma</code>	error standard deviation (for additive Gaussian errors).
<code>patchhw</code>	Half width of patches in each dimension (patchsize is $(2*\text{patchhw}+1)^d$ for d-dimensional array).
<code>searchhw</code>	Half width of search area (size of search area is $(2*\text{searchhw}+1)^d$ for d-dimensional array).
<code>pd</code>	If $\text{pd} < (2*\text{patchhw}+1)^d$ use pd principal components instead of complete patches.

Details

The implementation follows the description of the Non-Local-Means Filter of Buades et al 2005 on http://www.numerical-tours.com/matlab/denoisingadv_6_nl_means/#biblio that incorporates dimension reduction for patch comparisons by PCA.

Value

A list of class "nlmeans" with components

theta	Denoised array
lambda	Scale parameter used
sigma	The error standard deviation
patchhw	Half width of patches
pd	Effective patchsize used
searchhw	Half width of search area

Note

use `setCores='number of threads'` to enable parallel execution.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06 .

A. Buades, B. Coll and J. M. Morel (2006). A review of image denoising algorithms, with a new one. Simulation, 4, 490-530. DOI:10.1137/040616024.

http://www.numerical-tours.com/matlab/denoisingadv_6_nl_means/#biblio

paws

Adaptive weigths smoothing using patches

Description

The function implements a version the propagation separation approach that uses patches instead of individuel voxels for comparisons in parameter space. Functionality is analog to function [aws](#). Using patches allows for an improved handling of locally smooth functions and in 2D and 3D for improved smoothness of discontinuities at the expense of increased computing time.

Usage

```
paws(y, hmax = NULL, mask=NULL, onestep = FALSE, aws = TRUE, family = "Gaussian",
      lkern = "Triangle", aggkern = "Uniform", sigma2 = NULL, shape = NULL,
      scorr = 0, spmin = 0.25, ladjust = 1, wghts = NULL, u = NULL,
      graph = FALSE, demo = FALSE, patchsize = 1)
```

Arguments

y	array y containing the observe response (image intensity) data. dim(y) determines the dimensionality and extend of the grid design.
mask	logical array defining a mask. All computations are restricted to the mask.
hmax	hmax specifies the maximal bandwidth. Defaults to hmax=250, 12, 5 for 1D, 2D, 3D images, respectively. In case of lkern="Gaussian" the bandwidth is assumed to be given in full width half maximum (FWHM) units, i.e., 0.42466 times gridsize.
onestep	apply the last step only (use for test purposes only)
aws	logical: if TRUE structural adaptation (AWS) is used.
family	family specifies the probability distribution. Default is family="Gaussian", also implemented are "Bernoulli", "Poisson", "Exponential", "Volatility", "Variance" and "NCchi". family="Volatility" specifies a Gaussian distribution with expectation 0 and unknown variance. family="Volatility" specifies that $p*y/\theta$ is distributed as χ^2 with p=shape degrees of freedom. family="NCchi" uses a noncentral Chi distribution with p=shape degrees of freedom and noncentrality parameter theta
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.
aggkern	character: kernel used in stagewise aggregation, either "Triangle" or "Uniform"
sigma2	sigma2 allows to specify the variance in case of family="Gaussian". Not used if family!="Gaussian". Defaults to NULL. In this case a homoskedastic variance estimate is generated. If length(sigma2)==length(y) then sigma2 is assumed to contain the pointwise variance of y and a heteroscedastic variance model is used.
shape	Allows to specify an additional shape parameter for certain family models. Currently only used for family="Variance", that is χ -Square distributed observations with shape degrees of freedom.
scorr	The vector scorrr allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).
spmin	Determines the form (size of the plateau) in the adaptation kernel. Not to be changed by the user.
ladjust	factor to increase the default value of lambda
wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with u=0
graph	If graph=TRUE intermediate results are illustrated after each iteration step. Defaults to graph=FALSE.

demo	If demo=TRUE the function pauses after each iteration. Defaults to demo=FALSE.
patchsize	positive integer defining the size of patches. Number of grid points within the patch is $(2 \times \text{patchsize} + 1)^d$ with d denoting the dimensionality of the design.

Details

see [aws](#). The procedure is supposed to produce superior results if the assumption of a local constant image is violated or if smoothness of discontinuities is desired.

Value

returns an object of class `aws` with slots

```

y = "numeric"    y
dy = "numeric"   dim(y)
x = "numeric"    numeric(0)
ni = "integer"   integer(0)
mask = "logical" logical(0)
theta = "numeric" Estimates of regression function, length: length(y)
hseq = "numeric"  sequence of bandwidths employed
mae = "numeric"   Mean absolute error for each iteration step if u was specified, numeric(0) else
psnr = "numeric"  Peak signal-to-noise ratio for each iteration step if u was specified, numeric(0)
                  else
var = "numeric"   approx. variance of the estimates of the regression function. Please note that
                  this does not reflect variability due to randomness of weights.
xmin = "numeric"  numeric(0)
xmax = "numeric"  numeric(0)
wghts = "numeric" numeric(0), ratio of distances wghts[-1]/wghts[1]
degree = "integer" 0
hmax = "numeric"   effective hmax
sigma2 = "numeric" provided or estimated error variance
scorr = "numeric"  scorr
family = "character" family

```

```

shape = "numeric"
      shape
lkern = "integer"
      integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
      5="Gaussian"
lambda = "numeric"
      effective value of lambda
ladjust = "numeric"
      effective value of ladjust

aws = "logical"  aws
memory = "logical"
      memory
homogen = "logical"
      homogen
earlystop = "logical"
      FALSE
varmodel = "character"
      "Constant"
vcoef = "numeric"
      numeric(0)
call = "function"
      the arguments of the call to aws

```

Note

use `setCores='number of threads'` to enable parallel execution.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

J. Polzehl, K. Tabelow (2019). Magnetic Resonance Brain Imaging: Modeling and Data Analysis Using R. Springer, Use R! series. Appendix A. Doi:10.1007/978-3-030-29184-6.

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06 .

See Also

See also [aws](#), [lpaws](#), [vpaws](#), `link{awsdata}`

Examples

```

## Not run:
setCores(2)
y <- array(rnorm(64^3), c(64, 64, 64))
yhat <- paws(y, hmax=6)

```

```
## End(Not run)
```

plot-methods

Methods for Function 'plot' from package 'graphics' in Package 'aws'

Description

Visualization of objects of class "aws", "awssegment", "kernsm" and "ICIsMOOTH"

Methods

signature(x = "ANY") Generic function: see [plot](#).

signature(x = "aws") Visualization of objects of class "aws"

signature(x = "awssegment") Visualization of objects of class "awssegment"

signature(x = "ICIsMOOTH") Visualization of objects of class "ICIsMOOTH"

signature(x = "kernsm") Visualization of objects of class "kernsm"

Author(s)

Jörg Polzehl <polzehl@wias-berlin.de>

See Also

[aws](#), [awssegment](#), [ICIsMOOTH](#) [kernsm](#)

print-methods

Methods for Function 'print' from package 'base' in Package 'aws'

Description

The function provides information on data dimensions, creation of the object and existing slot-names for objects of class "aws", "awssegment", "ICIsMOOTH" and "kernsm"

Methods

signature(x = "ANY") Generic function: see [print](#).

signature(x = "aws") Provide information on data dimensions, creation of the object and existing slot-names for objects of class "aws"

signature(x = "awssegment") Provide information on data dimensions, creation of the object and existing slot-names for objects of class "awssegment"

signature(x = "ICIsMOOTH") Provide information on data dimensions, creation of the object and existing slot-names for objects of class "ICIsMOOTH"

signature(x = "kernsm") Provide information on data dimensions, creation of the object and existing slot-names for objects of class "kernsm"

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

See Also

[aws](#), [awssegment](#), [ICISmooth](#) [kernsm](#)

qmeasures

Quality assessment for image reconstructions.

Description

Computes selected criteria for quality assessments of

Usage

```
qmeasures(img, ref,  
          which = c("PSNR", "MAE", "MSE", "RMSE", "SSIM", "MAGE", "RMSGE"),  
          mask = FALSE)
```

Arguments

img	2D/3D image, object of class "aws", "ICISmooth", "kernsm", "nlmeans" or array.
ref	Reference image (array, matrix or vector) for comparison.
which	Criterion to use for Quality assessment. Please specify a subset of "PSNR" (Peak Signal to Noise Ratio), "MAE" (Mean Absolute Error), "MSE" (Mean Squared Error), "RMSE" (Root Mean Squared Error), "SSIM" (Structural SIMilarity), "MAGE" (Mean Absolute Gradient Error), "RMSGE" (Root Mean Squared Gradient Error).
mask	Logical of same dimension as img/ref. Calculation can be restricted to mask.

Details

Calculates specified quality indices.

Value

A vector with names as specified in which.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

Description

Methods function `risk` in package **aws**. For an given array `u` the following statistics are computed : Root Mean Squared Error $RMSE \leftarrow \sqrt{\text{mean}((y-u)^2)}$, Signal to Noise Ratio $SNR \leftarrow 10 \cdot \log(\text{mean}(u^2)/MSE, 10)$, Peak Signal to Noise Ratio $PSNR \leftarrow 10 \cdot \log(\max(u^2)/MSE, 10)$, Mean Absolute Error $MAE \leftarrow \text{mean}(\text{abs}(y-u))$, Maximal Absolute Error $MaxAE \leftarrow \max(\text{abs}(y-u))$, Universal Image Quality Index (UIQI) (Wang and Bovik (2002)).

Usage

```
## S4 method for signature 'array'
risk(y, u=0)
## S4 method for signature 'aws'
risk(y, u=0)
## S4 method for signature 'awssegment'
risk(y, u=0)
## S4 method for signature 'ICIsMOOTH'
risk(y, u=0)
## S4 method for signature 'kernsm'
risk(y, u=0)
## S4 method for signature 'numeric'
risk(y, u=0)
```

Arguments

<code>y</code>	object
<code>u</code>	array of dimension <code>dim(y)</code> or <code>dim(extract(y, what="yhat"))\$y</code> or scalar value used in comparisons.

Methods

`signature(y = "ANY")` The method extract and/or compute specified statistics from object of class

`signature(y = "array")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

`signature(y = "aws")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

`signature(y = "awssegment")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

`signature(y = "ICIsMOOTH")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

`signature(y = "kernsm")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

`signature(y = "numeric")` Returns a list with components `RMSE`, `SNR`, `PSNR`, `MAE`, `MaxAE`, `UIQI`

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>

References

V. Katkovnik, K. Egiazarian and J. Astola, *Local Approximation Techniques in Signal And Image Processing*, SPIE Society of Photo-Optical Instrumentation Engin., 2006, PM157

Z. Wang and A. C. Bovik, *A universal image quality index*, IEEE Signal Processing Letters, vol. 9, N3, pp. 81-84, 2002.

show-methods

Methods for Function 'show' in Package 'aws'

Description

The function provides information on data dimensions, data source and existing slot-names for objects of class "aws", "awssegment", "ICIsMOOTH" and "kernsm" in package **aws**

Methods

signature(object = "ANY") Generic function.

signature(object = "aws") Provide information on data dimensions, data source and existing slot-names for objects of class "dti" and classes that extent "aws".

signature(object = "awssegment") Provide information on data dimensions, data source and existing slot-names for objects of class "dti" and classes that extent "awssegment".

signature(object = "ICIsMOOTH") Provide information on data dimensions, data source and existing slot-names for objects of class "dti" and classes that extent "ICIsMOOTH".

signature(object = "kernsm") Provide information on data dimensions, data source and existing slot-names for objects of class "dti" and classes that extent "kernsm".

Author(s)

Karsten Tabelow <tabelow@wias-berlin.de>

Joerg Polzehl <polzehl@wias-berlin.de>

See Also

[aws](#), [awssegment](#), [ICIsMOOTH](#) [kernsm](#)

Description

smooth3D and medianFilter3D are auxiliary functions for non-adaptive smoothing of 3D image data using kernel or median smoothing. Both function restrict to sub-areas determined by a mask. The functions are used in packages dti and qMRI.

Functions aws3Dmask and aws3Dmaskfull perform adaptive weights smoothing on statistical parametric maps in fMRI. Variability of results is determined from smoothed (using the same weighting schemes) residuals in order to correctly account for spatial correlation. These functions are intended to be used internally in package fmri. They have been moved here because they share significant parts of the openMP parallelized Fortran code underlying function aws.

Usage

```
smooth3D(y, h, mask, lkern = "Gaussian", weighted = FALSE, sigma2 = NULL,
        wghts = NULL)
medianFilter3D(y, h = 10, mask = NULL)
aws3Dmask(y, mask, lambda, hmax, res = NULL, sigma2 = NULL, lkern = "Gaussian",
        skern = "Plateau", weighted = TRUE, u = NULL, wghts = NULL,
        h0 = c(0, 0, 0), testprop = FALSE)
aws3Dmaskfull(y, mask, lambda, hmax, res = NULL, sigma2 = NULL, lkern = "Gaussian",
        skern = "Plateau", weighted = TRUE, u = NULL, wghts = NULL,
        testprop = FALSE)
```

Arguments

y	3D array of data in case of functions smooth3D and medianFilter3D. For aws3Dmask* with !is.null(mask) a vector of length sum(mask) containing only data values within the specified mask.
lkern	character: location kernel, either "Triangle", "Plateau", "Quadratic", "Cubic" or "Gaussian". The default "Triangle" is equivalent to using an Epanechnikov kernel, "Quadratic" and "Cubic" refer to a Bi-weight and Tri-weight kernel, see Fan and Gijbels (1996). "Gaussian" is a truncated (compact support) Gaussian kernel. This is included for comparisons only and should be avoided due to its large computational costs.
weighted	logical: use inverse variances as weights.
sigma2	sigma2 allows to specify the variance of data entries.
mask	optional logical mask, same dimensionality as y
h	bandwidth to use. In case of lkern="Gaussian" this is in FWHM (full width half maximum) units. Value refers to first voxel dimension.
wghts	voxel dimensions. Defaults to c(1, 1, 1)
lambda	kritical scale parameter in hypothesis testing (adaptive weights smoothing)

hmax	maximum bandwidth for adaptive weights smoothing
res	array of residuals with dimension $c(nres, \text{sum}(\text{mask}))$.
skern	skern specifies the kernel for the statistical penalty. Defaults to "Plateau", the alternatives are "Triangle" and "Exp". "Plateau" specifies a kernel that is equal to 1 in the interval (0,.3), decays linearly in (.3,1) and is 0 for arguments larger than 1. lkern="Plateau" and lkern="Triangle" allow for much faster computation (saves up to 50% CPU-time). lkern="Plateau" produces a less random weighting scheme.
u	For test purposes in simulations: noiseless 3D data.
h0	Vector of 3 bandwidths corresponding to a Gaussian kernel that would produce a comparable spatial correlation by convoluting iid data.
testprop	logical: test the validity of a propagation condition for the specified value of lambda.

Value

Functions smooth3D and medianFilter3D return a 3D array. Functions awsmask* return a list with smoothed values of y in component theta and smoothed residuals in component res.

Note

Functions awsmask* are used intenally in package fmri. They refer to the situation, typical for fMRI, where the data are spatially correlated and this correlation can be accessed using residuals with respect to a model.

Author(s)

Joerg Polzehl <polzehl@wias-berlin.de>, Karsten Tabelow <tabelow@wias-berlin.de>

smse3ms

Adaptive smoothing in orientation space SE(3)

Description

The functions perform adaptive weights smoothing for data in orientation space SE(3), e.g. diffusion weighted MR data, with spatial coordinates given by voxel location within a mask and spherical information given by gradient direction. Observations can belong to different shells characterized by b-value bv. The data provided should only refer to voxel within mask.

Usage

```
smse3ms(sb, s0, bv, grad, kstar, lambda, kappa0, mask, sigma,
        ns0 = 1, ws0 = 1, vext = NULL, ncoils = 1, verbose = FALSE, usemaxni = TRUE)
smse3(sb, s0, bv, grad, mask, sigma, kstar, lambda, kappa0,
       ns0 = 1, vext = NULL, vred = 4, ncoils = 1, model = 0, dist = 1,
       verbose = FALSE)
```

Arguments

sb	2D array of diffusion weighted data, first dimension refers to index of voxel within the mask, second dimension to the number of diffusion weighted images.
s0	vector of length sum(mask) containing values within mask of an average non-diffusion-weighted image.
bv	vector of b-values.
grad	matrix of gradient directions with dim(grad)[1]==3.
kstar	number of steps in adaptive weights smoothing.
lambda	Scale parameter in adaptation
kappa0	determines amount of smoothing on the sphere. Larger values correspond to stronger smoothing on the sphere. If kappa0=NULL a value is that corresponds to a variance reduction with factor vred on the sphere.
mask	3D image defining a mask (logical)
sigma	Error standard deviation. Assumed to be known and homogeneous in the current implementation. A reasonable estimate may be defined as the modal value of standard deviations obtained using method getsdofsb.
ns0	Actual number of non-diffusion-weighted images used to obtain s0 by averaging.
ws0	Weight for non-diffusion-weighted images in statistical penalty.
vext	Voxel extensions.
ncoils	Effective number of receiver coils (in case of e.g. GRAPPA reconstructions), should be 1 in case of SENSE reconstructions. 2*ncoils is the number of degrees of freedom of the intensity distribution used.
verbose	If verbose=TRUE additional reports are given.
usemaxni	If "usemaxni==TRUE" a stricter penalization is used.
vred	Used if kappa0=NULL to specify the variance reduction on the sphere when suggesting a value of kappa0.
model	Determines which quantities are smoothed. Possible values are "Chi" for observed values (assumed to be distributed as noncentral Chi with 2*ncoils degrees of freedom), "Chi2" for squares of observed values (assumed to be distributed as noncentral Chi-squared with 2*ncoils degrees of freedom). "Gapprox" and "Gapprox2" use a Gaussian approximation for the noncentral Chi distribution to smooth observed and squared values, respectively.
dist	Distance in SE3. Reasonable values are 1 (default, see Becker et.al. 2012), 2 (a slight modification of 1: with $k6^2$ instead of $\text{abs}(k6)$) and 3 (using a 'naive' distance on the sphere)

Value

The functions return lists with main results in components th and th0 containing the smoothed data.

Note

These functions are intended to be used internally in package `dti` only.

Author(s)

Jörg Polzehl <polzehl@wias-berlin.de>

References

Joerg Polzehl, Karsten Tabelow (2019). Magnetic Resonance Brain Imaging: Modeling and Data Analysis Using R. Springer, Use R! series. Doi:10.1007/978-3-030-29184-6.

S. Becker, K. Tabelow, H.U. Voss, A. Anwender, R. Heidemann, J. Polzehl. Position-orientation adaptive smoothing of diffusion weighted magnetic resonance data (POAS). *Medical Image Analysis*, 2012, 16, 1142-1155. DOI:10.1016/j.media.2012.05.007.

S. Becker, K. Tabelow, S. Mohammadi, N. Weiskopf, J. Polzehl. Adaptive smoothing of multi-shell diffusion-weighted magnetic resonance data by msPOAS. *Neuroimage*, 2014, 95, 90-105. DOI:10.1016/j.neuroimage.2014.03.053.

summary-methods	<i>Methods for Function 'summary' from package 'base' in Package 'aws'</i>
-----------------	--

Description

The method provides summary information for objects of class `"aws"`, `"awssegment"`, `"ICIsmooth"` and `"kernsm"`

Methods

`signature(object = "ANY")` Generic function: see [summary](#).

`signature(object = "aws")` The function provides summary information for objects of class `"aws"`

`signature(object = "awssegment")` The function provides summary information for objects of class `"awssegment"`

`signature(object = "ICIsmooth")` The function provides summary information for objects of class `"ICIsmooth"`

`signature(object = "kernsm")` The function provides summary information for objects of class `"kernsm"`

Author(s)

Jörg Polzehl <polzehl@wias-berlin.de>

See Also

[aws](#), [awssegment](#), [ICIsmooth](#) [kernsm](#)

TV_denoising	<i>TV/TGV denoising of image data</i>
--------------	---------------------------------------

Description

Total variation and total generalized variation are classical energy minimizing methods for image denoising.

Usage

```
TV_denoising(datanoisy, alpha, iter = 1000, tolmean = 1e-06,
             tolsup = 1e-04, scale = 1, verbose=FALSE)
TGV_denoising(datanoisy, alpha, beta, iter = 1000, tolmean = 1e-06,
              tolsup = 1e-04, scale = 1, verbose=FALSE)
TV_denoising_colour(datanoisy, alpha, iter = 1000, tolmean = 1e-06,
                   tolsup = 1e-04, scale = 1, verbose=FALSE)
TGV_denoising_colour(datanoisy, alpha, beta, iter = 1000, tolmean = 1e-06,
                    tolsup = 1e-04, scale = 1, verbose=FALSE)
```

Arguments

datanoisy	matrix of noisy 2D image data. In case of TV_denoising_colour and TGV_denoising_colour and array with third dimension referring to RGB channels.
alpha	TV regularization parameter.
beta	additional TGV regularization parameter.
iter	max. number of iterations
tolmean	requested accuracy for mean image correction
tolsup	requested accuracy for max (over pixel) image correction
scale	image scale
verbose	report convergence diagnostics.

Details

Reimplementation of original matlab code by Kostas Papafitsoros (WIAS).

Value

TV/TGV reconstructed image data (2D array)

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

- J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06.
- Rudin, L.I., Osher, S. and Fatemi, E. (1992). Nonlinear total variation based noise removal algorithms. Phys. D, 60, 259-268. DOI: 10.1016/0167-2789(92)90242-F.
- Bredies, K., Kunisch, K. and Pock, T. (2010). Total Generalized Variation. SIAM J. Imaging Sci., 3, 492-526. DOI:10.1137/090769521.

vaws	<i>vector valued version of function aws The function implements the propagation separation approach to nonparametric smoothing (formerly introduced as Adaptive weights smoothing) for varying coefficient likelihood models with vector valued response on a 1D, 2D or 3D grid.</i>
------	---

Description

The function implements a version the propagation separation approach that uses vector valued instead of scalar responses.

Usage

```
vaws(y, kstar = 16, sigma2 = 1, mask = NULL, scorr = 0, spmin = 0.25,
      ladjust = 1, wghts = NULL, u = NULL, maxni = FALSE)
vawscov(y, kstar = 16, invcov = NULL, mask = NULL, scorr = 0, spmin = 0.25,
        ladjust = 1, wghts = NULL, u = NULL, maxni = FALSE)
```

Arguments

y	y contains the observed response data. <code>dim(y)</code> determines the dimensionality and extend of the grid design. First component varies over components of the response vector.
kstar	maximal number of steps to employ. Determines maximal bandwidth.
sigma2	specifies a homogeneous error variance.
invcov	array of voxelwise inverse covariance matrixes, first index corresponds to upper diagonal inverse covariance matrix.
mask	logical mask. All computations are restrikted to design poins within the mask.
scorr	The vector <code>scorr</code> allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).
spmin	determines the form (size of the plateau) in the adaptation kernel. Not to be changed by the user.
ladjust	factor to increase the default value of lambda

wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with $u=0$
maxni	If TRUE use $\max_{l \leq k} (N_i^{(l)})$ instead of $(N_i^{(k)})$ in the definition of the statistical penalty.

Details

see [aws](#). Expets vector valued responses. Currently only implements the case of additive Gaussian errors.

Value

returns an object of class `aws` with slots

<code>y = "numeric"</code>	<code>y</code>
<code>dy = "numeric"</code>	<code>dim(y)</code>
<code>x = "numeric"</code>	<code>numeric(0)</code>
<code>ni = "integer"</code>	<code>integer(0)</code>
<code>mask = "logical"</code>	<code>logical(0)</code>
<code>theta = "numeric"</code>	Estimates of regression function, length: <code>length(y)</code>
<code>hseq = "numeric"</code>	sequence of bandwidths employed
<code>mae = "numeric"</code>	Mean absolute error for each iteration step if <code>u</code> was specified, <code>numeric(0)</code> else
<code>psnr = "numeric"</code>	Peak signal-to-noise ratio for each iteration step if <code>u</code> was specified, <code>numeric(0)</code> else
<code>var = "numeric"</code>	approx. variance of the estimates of the regression function. Please note that this does not reflect variability due to randomness of weights.
<code>xmin = "numeric"</code>	<code>numeric(0)</code>
<code>xmax = "numeric"</code>	<code>numeric(0)</code>
<code>wghts = "numeric"</code>	<code>numeric(0)</code> , ratio of distances <code>wghts[-1]/wghts[1]</code>
<code>degree = "integer"</code>	0
<code>hmax = "numeric"</code>	effective <code>hmax</code>
<code>sigma2 = "numeric"</code>	provided or estimated (inverse) error variance

```

scorr = "numeric"
      scorrr
family = "character"
      family
shape = "numeric"
      shape
lkern = "integer"
      integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
      5="Gaussian"
lambda = "numeric"
      effective value of lambda
ladjust = "numeric"
      effective value of ladjust

aws = "logical"  aws
memory = "logical"
      memory
homogen = "logical"
      homogen
earlystop = "logical"
      FALSE
varmodel = "character"
      "Constant"
vcoef = "numeric"
      numeric(0)
call = "function"
      the arguments of the call to aws

```

Note

use `setCores='number of threads'` to enable parallel execution.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

- J. Polzehl, K. Tabelow (2019). Magnetic Resonance Brain Imaging: Modeling and Data Analysis Using R. Springer, Use R! series. Appendix A. Doi:10.1007/978-3-030-29184-6.
- J. Polzehl, V. Spokoiny, Adaptive Weights Smoothing with applications to image restoration, J. R. Stat. Soc. Ser. B Stat. Methodol. 62 , (2000) , pp. 335–354. DOI:10.1111/1467-9868.00235.
- J. Polzehl, V. Spokoiny, Propagation-separation approach for local likelihood estimation, Probab. Theory Related Fields 135 (3), (2006) , pp. 335–362. DOI:10.1007/s00440-005-0464-1.

See Also

See also [aws](#), [vpaws](#), `link{awsdata}`

Examples

```
## Not run:
setCores(2)
y <- array(rnorm(4*64^3),c(4,64,64,64))
yhat <- vaws(y,kstar=20)

## End(Not run)
```

vpaws	<i>vector valued version of function paws with homogeneous covariance structure</i>
-------	---

Description

The function implements a vector-valued version the propagation separation approach that uses patches instead of individual voxels for comparisons in parameter space. Functionality is analog to function [vaws](#). Using patches allows for an improved handling of locally smooth functions and in 2D and 3D for improved smoothness of discontinuities at the expense of increased computing time.

Usage

```
vpaws(y, kstar = 16, sigma2 = 1, invcov = NULL, mask = NULL, scor = 0, spmin = 0.25,
      ladjust = 1, wghts = NULL, u = NULL, patchsize = 1)
vpawscov(y, kstar = 16, invcov = NULL, mask = NULL, scor = 0, spmin = 0.25, ladjust = 1,
          wghts = NULL, maxni = FALSE, patchsize = 1)
vpawscov2(y, kstar = 16, invcov = NULL, mask = NULL, scor = 0, spmin = 0.25,
           lambda = NULL, ladjust = 1, wghts = NULL, patchsize = 1,
           data = NULL, verbose = TRUE)
```

Arguments

y	y can be a full array of vector valued data, or, if mask is provided, be a matrix with columns corresponding to points/pixel/voxel within the mask. In the first case <code>dim(y)</code> determines the dimensionality and extend of the grid design, in the second case this information is obtained from the dimensions of mask. the first component varies over components of the response vector.
kstar	maximal number of steps to employ. Determines maximal bandwidth.
sigma2	specifies a homogeneous error variance.
invcov	array (or matrix) of voxelwise inverse covariance matrixes, first index corresponds to upper diagonal inverse covariance matrix.
mask	logical mask. All computations are restricted to design points within the mask.
scor	The vector <code>scor</code> allows to specify a first order correlations of the noise for each coordinate direction, defaults to 0 (no correlation).
spmin	determines the form (size of the plateau) in the adaptation kernel. Not to be changed by the user.

ladjust	factor to increase the default value of lambda
wghts	wghts specifies the diagonal elements of a weight matrix to adjust for different distances between grid-points in different coordinate directions, i.e. allows to define a more appropriate metric in the design space.
u	a "true" value of the regression function, may be provided to report risks at each iteration. This can be used to test the propagation condition with $u=0$
patchsize	positive integer defining the size of patches. Number of grid points within the patch is $(2*patchsize+1)^d$ with d denoting the dimensionality of the design.
maxni	require growing sum of weights
lambda	explicit value of lambda
data	optional vector-valued images to be smoothed using the weighting scheme of the last step
verbose	logical: provide information on progress.

Details

see [vaws](#). Parameter y The procedure is supposed to produce superior results if the assumption of a local constant image is violated or if smoothness of discontinuities is desired.

Function `vpawscov2` is intended for internal use in package `qMRI` only.

Value

function `vpaws` returns returns an object of class `aws` with slots

<code>y = "numeric"</code>	<code>y</code>
<code>dy = "numeric"</code>	<code>dim(y)</code>
<code>x = "numeric"</code>	<code>numeric(0)</code>
<code>ni = "integer"</code>	<code>integer(0)</code>
<code>mask = "logical"</code>	<code>logical(0)</code>
<code>theta = "numeric"</code>	Estimates of regression function, length: <code>length(y)</code>
<code>hseq = "numeric"</code>	sequence of bandwidths employed
<code>mae = "numeric"</code>	Mean absolute error for each iteration step if <code>u</code> was specified, <code>numeric(0)</code> else
<code>psnr = "numeric"</code>	Peak signal-to-noise ratio for each iteration step if <code>u</code> was specified, <code>numeric(0)</code> else
<code>var = "numeric"</code>	approx. variance of the estimates of the regression function. Please note that this does not reflect variability due to randomness of weights. Currently also uses factor $1/ni$ instead of the correct $\sum(w_{ij}^2)/ni^2$
<code>xmin = "numeric"</code>	<code>numeric(0)</code>

```

xmax = "numeric"
        numeric(0)
wghts = "numeric"
        numeric(0), ratio of distances wghts[-1]/wghts[1]
degree = "integer"
        0
hmax = "numeric"
        effective hmax
sigma2 = "numeric"
        provided or estimated error variance
scorr = "numeric"
        scor
family = "character"
        family
shape = "numeric"
        shape
lkern = "integer"
        integer code for lkern, 1="Plateau", 2="Triangle", 3="Quadratic", 4="Cubic",
        5="Gaussian"
lambda = "numeric"
        effective value of lambda
ladjust = "numeric"
        effective value of ladjust

aws = "logical"  aws
memory = "logical"
        memory
homogen = "logical"
        homogen
earlystop = "logical"
        FALSE
varmodel = "character"
        "Constant"
vcoef = "numeric"
        numeric(0)
call = "function"
        the arguments of the call to aws

```

If `y` contained only information (condensed data) for positions within a mask, then the returned object only contains results for these positions.

Note

use `setCores='number of threads'` to enable parallel execution.

Author(s)

Joerg Polzehl, <polzehl@wias-berlin.de>, <https://www.wias-berlin.de/people/polzehl/>

References

J. Polzehl, K. Tabelow (2019). Magnetic Resonance Brain Imaging: Modeling and Data Analysis Using R. Springer, Use R! series. Appendix A. Doi:10.1007/978-3-030-29184-6.

J. Polzehl, K. Papafitsoros, K. Tabelow (2020). Patch-Wise Adaptive Weights Smoothing in R, Journal of Statistical Software, 95(6), 1-27. doi:10.18637/jss.v095.i06 .

See Also

See also [vaws](#), [lpaws](#), [vawscov](#), [link{awsdata}](#)

Examples

```
## Not run:
setCores(2)
y <- array(rnorm(4*64^3),c(4,64,64,64))
yhat <- vpaws(y,kstar=20)

## End(Not run)
```

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