

Package ‘AdaptFitOS’

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Title Adaptive Semiparametric Regression with Simultaneous Confidence Bands

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Depends SemiPar, nlme, MASS, mgcv

Suggests splines

Description Based on the function ``asp" of the AdaptFit package fits semiparametric regression models with spatially adaptive penalized splines and computes simultaneous confidence bands.

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AdaptFitOS-package	<i>Adaptive Semiparametric Regression with Simultaneous Confidence Bands</i>
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Description

Based on package `AdaptFit` (which is itself based on the `SemiPar` package) fits semiparametric regression models with spatially adaptive penalized splines and computes simultaneous confidence bands.

Particular differences to `AdaptFit` include the availability of simultaneous confidence bands and B-spline basis functions and different functionality of the `plot` function. However, random effects, autocorrelations and interaction surfaces are not supported. Further, only Gaussian responses are supported. Note that in contrast to `AdaptFit` and `SemiPar`, estimated curves are centered to have zero mean. Furthermore, unlike in package `SemiPar`, dummies for categorical covariates are constructed automatically if a variable is given as factor. and unlike `SemiPar`, categorical covariates are automatically detected. For computation of the critical value for simultaneous confidence bands based on Hotelling's volume-of-tube formula, some functions of the `libtube` library by Catherine Loader (see package `locfit`) are used. See the references for details on the construction of the confidence bands.

Details

```
Package:  AdaptFitOS
Version:  0.42
Date:    2012-06-03
Depends: SemiPar, MASS, nlme, cluster
```

Index:

<code>asp2</code>	Fit a semiparametric regression model with spatially adaptive penalized
<code>aspFormula</code>	An asp formula
<code>aspHetero</code>	Estimate varying residual variance
<code>fitted.asp</code>	Fitted values for semiparametric regression.
<code>plot.asp</code>	Plots fitted curves or their derivatives including simultaneous confidence bands
<code>plot.scbm</code>	Plots fitted curves in a <code>scbm</code> object including simultaneous confidence bands
<code>predict.asp</code>	Semiparametric regression prediction.
<code>residuals.asp</code>	Residuals for semiparametric regression.
<code>scbm</code>	Calculate simultaneous confidence bands for penalized splines
<code>summary.asp</code>	Semiparametric regression summary

The function `asp2()` is used to fit the model. Using the resulting `asp` object, fitted curves or their derivatives can be plotted with `plot.asp` and information on the parametric effects can be printed using `summary`.

Author(s)

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References

- Ganguli, B. and Wand, M.P. (2005)
SemiPar 1.0 Users' Manual.
<http://www.uow.edu.au/~mwand/SPmanu.pdf>
- Krivobokova, T., Crainiceanu, C.M. and Kauermann, G. (2008)
Fast Adaptive Penalized Splines. *Journal of Computational and Graphical Statistics*, 17(1):1-20.
- Krivobokova, T., Kneib, T., and Claeskens, G. (2010)
Simultaneous confidence bands for penalized spline estimators. *Journal of the American Statistical Association*, 105(490):852-863.
- Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition. *Journal of the American Statistical Association*, 107(500): 1286-1296.

See Also

`spm` (package SemiPar), `asp` (package AdaptFit)

asp2	<i>Fit a semiparametric regression model with spatially adaptive penalized splines</i>
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Description

Fits semiparametric regression models using the mixed model representation of penalized splines with spatially adaptive penalties based on the `asp` function of the AdaptFit-package (which is based on `spm` function of SemiPar package). Particular differences to AdaptFit include the availability of simultaneous confidence bands and B-spline basis functions and different functionality of the `plot` function. However, random effects, autocorrelations and interaction surfaces are not supported. Further, only Gaussian responses are supported. Note that in contrast to AdaptFit and SemiPar, estimated curves are centered to have zero mean.

Usage

```
asp2(form, spar.method = "REML", contrasts=NULL,  
      omit.missing = NULL, returnFit=FALSE,  
      niter = 20, niter.var = 50, tol=1e-6, tol.theta=1e-6,  
      control=NULL)
```

Arguments

<code>form</code>	a formula describing the model to be fitted. See aspFormula for further information. Note, that an intercept is always included, whether given in the formula or not.
<code>spar.method</code>	method for automatic smoothing parameter selection. May be "REML" (restricted maximum likelihood) or "ML" (maximum likelihood).
<code>contrasts</code>	an optional list. See the <code>contrasts.arg</code> of model.matrix.default .
<code>omit.missing</code>	a logical value indicating whether fields with missing values are to be omitted.
<code>niter</code>	a maximum number of iterations for the mean estimation, default is 20.
<code>niter.var</code>	a maximum number of iterations for the variance of random effects estimation, default is 50.
<code>tol</code>	tolerance for the convergence criterion. Default is 1e-6.
<code>tol.theta</code>	tolerance for the convergence criterion (smoothing parameter function routine). Default is 1e-6.
<code>returnFit</code>	a logical value indicating whether the fitted object should be returned when the maximum number of iterations is reached without convergence of the algorithm. Default is FALSE.
<code>control</code>	see <code>lmeControl</code> in the documentation to <code>nlme</code> .

Value

A list object of class `asp` containing the fitted model. The components are:

<code>fitted</code>	fitted values.
<code>coef.mean</code>	estimated mean coefficients.
<code>design.matrices</code>	design matrices both for knots und subknots.
<code>x</code>	x values.
<code>knots</code>	knots.
<code>y.cov</code>	estimated covariance matrix of the response.
<code>random.var</code>	estimated covariance matrix of the random effects.
<code>subknots</code>	subknots.
<code>coef.random</code>	estimated spline coefficients of the covariance matrix of the random effects.
<code>var.random.var</code>	estimated variance of the spline coefficients of the covariance matrix of the random effects.
<code>fit</code>	mimics fit object of <code>lme()</code> .
<code>info</code>	information about the inputs.
<code>aux</code>	auxiliary information such as variability estimates.

Author(s)

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References

- Krivobokova, T., Crainiceanu, C.M. and Kauermann, G. (2008)
Fast Adaptive Penalized Splines. *Journal of Computational and Graphical Statistics*. 17(1) 1-20.
- Ruppert, D., Wand, M.P. and Carroll, R.J. (2003)
Semiparametric Regression Cambridge University Press.
<http://stat.tamu.edu/~carroll/semiregbook/>
- Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition.
Journal of the American Statistical Association, 107(500): 1286-1296.

See Also

[gam](#) (in package 'mgcv'), [asp](#) (in package 'AdaptFit'), [lme](#) (in package 'nlme')

Examples

```
## Not run:
#####
# Examples as in package AdaptFit
## scatterplot smoothing
x <- 1:1000/1000
mu <- exp(-400*(x-0.6)^2)+
5*exp(-500*(x-0.75)^2)/3+2*exp(-500*(x-0.9)^2)
y <- mu+0.5*rnorm(1000)

#fit with default knots
y.fit <- asp2(y~f(x,adap=TRUE))
plot(y.fit,residuals=TRUE,lwd=2,scb.lwd=2,scb.lty="dashed")
# with shaded confidence region.
# Use scb.lty="blank" to plot the shades only.
plot(y.fit,residuals=TRUE,shade=TRUE,scb.lty="blank")

## one more scatterplot smoothing with specified knots and subknots
x <- 1:400/400
mu <- sqrt(x*(1-x))*sin((2*pi*(1+2^((9-4*6)/5)))/(x+2^((9-4*6)/5)))
y <- mu+0.2*rnorm(400)

kn <- default.knots(x,80)
kn.var <- default.knots(kn,20)

y.fit <- asp2(y~f(x,knots=kn))
y.fit2 <- asp2(y~f(x,knots=kn,var.knots=kn.var,adap=TRUE))
op <- par(mfrow = c(1, 2))
plot(y.fit)
plot(y.fit2)
par(op)

## additive models
x1 <- 1:300/300
x2 <- runif(300)
```

```

mu1 <- exp(-400*(x1-0.6)^2)+
  5*exp(-500*(x1-0.75)^2)/3+2*exp(-500*(x1-0.9)^2)
mu2 <- sin(2*pi*x2)
y2 <- mu1+mu2+0.3*rnorm(300)

y2.fit <- asp2(y2~f(x1,adap=TRUE)+f(x2,adap=TRUE))
# switch off adaptive fitting for the first function
y21.fit <- asp2(y2~f(x1,adap=FALSE)+f(x2,adap=TRUE))
op <- par(mfrow = c(2, 2))
plot(y2.fit)
plot(y21.fit)
par(op)

# Use predict to avoid centering of smooths in case of scatterplot
# smoothing
library(SemiPar)
data(fossil)
attach(fossil)
fit <- asp2(strontium.ratio~f(age,basis="tps",adap=FALSE))
newdata.age <- data.frame(age=seq(90,130,length.out=50) )
preds <- predict(fit,newdata=newdata.age,se=FALSE)
plot(age,strontium.ratio)
lines(newdata.age$age,preds,col="red")

#####
#more examples
beta=function(l,m,x)
  return(gamma(l+m)*(gamma(l)*gamma(m))^(1-x)^l*(1-x)^(m-1))
f1 = function(x) return((0.6*beta(30,17,x)+0.4*beta(3,11,x))*1/0.958)
f2 = function(z) return((sin(2*pi*(z-0.5))^2)*1/.3535)
f3 = function(z)
  return((exp(-400*(z-0.6)^2)+
    5/3*exp(-500*(z-0.75)^2)+2*exp(-500*(z-0.9)^2))*1/0.549)

set.seed(1)
N <- 500
x1 = runif(N,0,1)
x2 = runif(N,0,1)
x3 = runif(N,0,1)

kn1 <- default.knots(x1,40)
kn2 <- default.knots(x2,40)
kn3 <- default.knots(x3,40)
kn.var3 <- default.knots(kn3,5)

y <- f1(x1)+f2(x2)+f3(x3)+0.3*rnorm(N)

# semiparametric model
fit1= asp2(y~x1+f(x2,basis="os",degree=3,knots=kn2,adap=FALSE)
  +f(x3,basis="os",degree=3,
    knots=kn3,var.knots=kn.var3,adap=FALSE),

```

```

        niter = 20, niter.var = 200)
summary(fit1)
plot(fit1,pages=1)

# all effects flexible
# fit model with all smoothing parameters constant
fit2a= asp2(y~f(x1,basis="os",degree=3,knots=kn1,adap=FALSE)
           +f(x2,basis="os",degree=3,knots=kn2,adap=FALSE)
           +f(x3,basis="os",degree=3,knots=kn3,adap=FALSE),
           niter = 20, niter.var = 200)
plot(fit2a,pages=1)

# fit model with last smoothing parameter adaptive
fit2b= asp2(y~f(x1,basis="os",degree=3,knots=kn1,adap=FALSE)
           +f(x2,basis="os",degree=3,knots=kn2,adap=FALSE)
           +f(x3,basis="os",degree=3,knots=kn3,adap=TRUE,
             var.knots=kn.var3,var.basis="os",var.degree=3),
           niter = 20, niter.var = 200)

# plot smoothing parameter function for covariate x3.
# Note that in the case of B-splines additional knots are added,
# see references.
plot(seq(0,1,length.out=42), fit2b$y.cov/fit2b$random.var[85:126],
     ylab=expression(lambda(x3)),xlab="x3",type="l",lwd=3)

# compute 95
# You could skip this and use "fit2b" instead of "scb2b" later on, however,
# if N is large, computing the SCBs various times can take some time
# if you don't need fitted values and bounds for all covariate points
# (can be computationally intensive due to large matrix dimensions),
# set calc.stdev=F such that these are not computed.
scb2b<- scbM(fit2b,calc.stdev=FALSE)
plot(scb2b,pages=1)

# plot only f(x2).
plot(scb2b,select=2,mfrow=c(1,1),lwd=3,ylab="f(x2)",xlab="x2")
# plot.scbm (and plot.asp) returns fitted values and confidence limits,
# if you only need the returned object set plot=FALSE
pscb=plot(scb2b,plot=FALSE)
# add pointwise confidence intervals to the plot
polygon(c(pscb$grid.x[[2]], rev(pscb$grid.x[[2]])),
       c(pscb$fitted[[2]]+1.96*pscb$Stdev[[2]],
         rev(pscb$fitted[[2]]-1.96*pscb$Stdev[[2]])),
       col = grey(0.85), border = NA)
lines(pscb$grid.x[[2]],pscb$lcb[[2]],lty="dotted",lwd=3)
lines(pscb$grid.x[[2]],pscb$fitted[[2]],lwd=3)
lines(pscb$grid.x[[2]],pscb$ucb[[2]],lty="dotted",lwd=3)

# plot first derivative of f(x1)
scb2bdrv<- scbM(fit2b,drv=1,calc.stdev=FALSE)
plot(scb2bdrv,select=1)
#the following would give the same result

```

```

#x11();plot(fit2b,select=1,drv=1)
# different style
plot(scb2bdrv,select=1,scb.lty="blank",
     shade=TRUE,shade.col="steelblue")

## End(Not run)

```

aspFormula

An asp formula

Description

A formula to be used in [asp2](#). The formula is close to the one used in [asp](#) of package `AdaptFit`.

Unlike in package `SemiPar`, dummies for categorical covariates are constructed automatically if a variable is given as factor (with contrasts as set by `options("contrasts")` or specified by a list in argument `contrasts`). Note that only parametric interactions are supported and that interacting covariates have to be multiplied beforehand and given as a new variable in the formula. Smooth terms are given by

```
f(x,basis="os",degree=3,knots,var.knots,var.basis,var.degree,adap=TRUE)
```

with the following arguments:

Arguments

<code>x</code>	the covariate
<code>basis</code>	the spline basis function to be used. "trunc.poly" for truncated polynomials, "tps" for thin plate splines and "os" for B-splines. The use of B-splines is recommended. Note that in contrast to packages <code>SemiPar</code> and <code>AdaptFit</code> , "os" is the default.
<code>degree</code>	the degree of the basis. In the case of B-splines also a vector of the form $c(p,q)$ with p the B-spline degree and q the penalty order (the integrated q -th squared derivative is penalized, see references). If only a scalar is given q is chosen such that $p=2*q-1$. Defaults are <code>degree=3</code> (<code>basis="tps"</code>), <code>degree=1</code> (<code>basis="trunc.poly"</code>) and <code>degree=c(3,2)</code> (<code>basis="os"</code>), respectively.
<code>knots</code>	the knots to be used. Using e.g. <code>kn=default.knots(x,40)</code> beforehand leads to 40 quantile based knots in the case of "tps" and "trunc.poly" bases. In the case of B-splines ("os"), knots are always equidistant and are automatically generated with the number equal to the length of the vector of knots given plus boundary knots. If no knots are given the number of knots is automatically chosen to be equal to $\text{floor}(n/\max(4, \text{floor}(n/35))) - 1$.
<code>adap</code>	TRUE for spatially adaptive smoothing parameter
<code>var.knots</code>	the knots for the spline basis for adaptively estimating the smoothing parameter. Note that in package <code>AdaptFit</code> "var.knot" (i.e. without "s") is used instead. If missing the number of knots is automatically chosen to be equal to $\text{floor}(\text{knots}/\max(4, \text{floor}(\text{knots}/35))) - 1$.

var.basis	spline basis function for adaptive smoothing parameter estimation. If missing, the same basis as for estimation of f is used.
var.degree	spline degree for adaptive smoothing parameter estimation. If missing, the same degree as for estimation of f is used.
spar	the smoothing parameter if desired. Usually this is left unspecified, such that the smoothing parameter is estimated by restricted maximum likelihood (see references). Currently doesn't work for basis="os".
df	the number of degrees of freedom corresponding to the REML choice of smoothing parameter if desired. Usually this is left unspecified, such that the smoothing parameter is estimated by restricted maximum likelihood (see references). Currently doesn't work for basis="os".

aspHetero

Estimate varying residual variance

Description

Estimates a varying residual variance on basis of an asp object.

Usage

```
aspHetero(object, xx, nknots=5, knots=NULL, basis="os",
          degree=c(3,2), tol=1e-8, niter=100, niter.var=250)
```

Arguments

object	an asp object.
xx	the covariate.
nknots	the number of knots. Does not apply when knots are given.
knots	the knots. Does not apply if basis=="os". Otherwise, if NULL nknots equidistant knots are used.
basis	the spline basis: "os" (default), "trunc.poly" or "tps".
degree	the spline degree (and penalty order in case of B-splines). Defaults to c(3,2).
tol	tolerance for the convergence criterion. Default is 1e-8.
niter	a maximum number of iterations for residual variance function estimation, default is 100.
niter.var	a maximum number of iterations for the variance of random effects estimation within the residual variance function estimation routine, default is 250.

Value

An object of class asp with varying variances, with additional element `sigmax` including information on the spline of the varying variance.

Author(s)

Manuel Wiesenfarth <m.wiesenfarth at dkfz de>

References

Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition.
Journal of the American Statistical Association, 107(500): 1286-1296.

Examples

```
## Not run:
attach(mcycle)

y=accel
kn1 <- default.knots(times,20)
# fit model with constant residual variance
fit= asp2(accel~f(times,basis="os",degree=3,knots=kn1,adap=FALSE),
  niter = 20, niter.var = 200)

# fit model with varying residual variance
fith=aspHetero(fit,times,tol=1e-8)
op <- par(mfrow = c(1,3))
plot(fit);plot(fith)
#sigma() returns the fitted varying residual variance
plot(sort(times),sigma(fith)[order(times)],type="l")
par(op)

## End(Not run)
```

fitted.asp

Fitted values for semiparametric regression.

Description

Extracts fitted values from a semiparametric regression fit object.

Usage

```
## S3 method for class 'asp'
fitted(object,...)
```

Arguments

object a fitted asp object as produced by asp2().
... other possible arguments.

Details

Extracts fitted from a semiparametric regression fit object. The fitted are defined to be the set of values obtained when the predictor variable data are substituted into the fitted regression model.

Value

The vector of fitted.

See Also

[plot.asp](#), [predict.asp](#), [summary.asp](#), [residuals.asp](#), [asp](#) (package AdaptFit)

Examples

```
## Not run:
library(SemiPar)
data(fossil)
attach(fossil)
fit <- asp2(strontium.ratio~f(age))
plot(fit, bands=FALSE)
points(age,fitted(fit)-fit$coef[1],col="red")

## End(Not run)
```

plot.asp

Plots fitted curves or their derivatives

Description

Plots fitted curves or their derivatives together with simultaneous confidence bands.

Usage

```
## S3 method for class 'asp'
plot(x, select=NULL, drv=0, bands=TRUE, level=0.95, grid=50, pages=0,
plot=TRUE, ylim=NULL, xlab=NULL, ylab=NULL,
      scb.lwd=1, scb.lty="dotted", shade=FALSE, shade.col=grey(0.85),
      residuals=FALSE, residuals.col="steelblue",
      bayes=FALSE, rug=TRUE,...)
```

Arguments

x	an asp object
select	vector specifying which curves in an additive model should be plotted. If NULL, all curves are plotted.
drv	the derivative order. Defaults to 0, i.e. the estimated curves themselves are plotted. First and second derivatives are supported. Does not apply to objects created by scbM .

bands	TRUE in order to include simultaneous confidence bands.
grid	number of points used for the plot, default value 50.
plot	if FALSE no plot is given
yylim	vector or list of vecotrs of limits on y axes. If NULL limits are automatically chosen. If multiple curves are plotted and a two-dimensional vector is given, y axes for all curves will be equal. A list with length equal to the number of smooth curves in the model can be given to specify different y-axes for each smooth.
pages	The number of pages over which to spread the output as in package <code>mgcv</code> . For example, if <code>pages=1</code> then all terms will be plotted on one page in an automatic fashion. If <code>pages=0</code> (default) all graphics settings are left as they are.
level	the level of confidence (does not apply to objects created by <code>scbM</code>).
xlab	label for the x axis. A list with length equal to the number of smooth curves in the model can be given to specify different labels for each smooth.
ylab	label for the y axis. A list with length equal to the number of smooth curves in the model can be given to specify different labels for each smooth.
scb.lwd	line width for simultaneous confidence bands
scb.lty	line type for simultaneous confidence bands. Use <code>scb.lty="blank"</code> , if you only want to plot the shades.
shade	set to TRUE to produce shaded regions as simultaneous confidence bands for smooths
shade.col	define the color used for shading confidence bands
residuals	if TRUE, partial residuals are added to the plot
residuals.col	color of partial residuals
rug	adds a rug representation (1-d plot) of the data to the plot.
bayes	FALSE for simultaneous confidence bands with (approximate) frequentist coverage probability, TRUE for (approximate) Bayesian coverage probability. See Krivobokova et al. (2010) for details.
...	further arguments to be passed to <code>plot()</code>

Details

`plot.asp()` first calls `scbM` and then `plot.scbm()` to plot an `asp` object. If plotting takes long (because of a large data set) and you want to plot multiple times with different settings, use `scbM` and then plot the resulting `scbm` object with `plot.scbm()`. Different to packages `SemiPar` and `AdaptFit`, estimated fits are centred to have zero mean. The simultaneous confidence bands have (approximate) frequentist coverage probabilities with automatic bias correction (see references).

Value

<code>grid.x</code>	list of the grid values used
<code>fitted</code>	list of the fitted values on the grid
<code>lcb</code>	list of the lower bounds of the confidence bands

ucb	list of the upper bounds of the confidence bands
drv	the derivative order
Stdev.fit	the standard deviations on the grid
ylim	list of ylim used for plotting
residuals	the partial residuals.

Author(s)

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References

- Krivobokova, T., Kneib, T., and Claeskens, G. (2010)
 Simultaneous confidence bands for penalized spline estimators. *Journal of the American Statistical Association*, 105(490):852-863.
- Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
 Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition. *Journal of the American Statistical Association*, 107(500): 1286-1296.

See Also

[plot.spm](#) in package SemiPar

Examples

```
# see asp2()
```

predict.asp	<i>Semiparametric regression prediction.</i>
-------------	--

Description

Takes a fitted asp object produced by asp2 and obtains predictions at new data values.

Usage

```
## S3 method for class 'asp'
predict(object,newdata,se,...)
```

Arguments

object	a fitted asp object as produced by asp2().
newdata	a data frame containing the values of the predictors at which predictions are required. The columns should have the same name as the predictors. Further, minima and maxima should currently coincide with those of the predictors.
se	when this is TRUE standard error estimates are returned for each prediction. The default is FALSE.
...	other arguments.

Details

Takes a fitted asp object produced by `asp2()` and obtains predictions at new data values as specified by the 'newdata' argument. If 'se=TRUE' then standard error estimates are also obtained.

Value

If `se=FALSE` then a vector of predictions at 'newdata' is returned. If `se=TRUE` then a list with components named 'fit' and 'se' is returned. The 'fit' component contains the predictions. The 'se' component contains standard error estimates.

Author(s)

Manuel Wiesenfarth, based on implementation of M.P. Wand (package SemiPar).

See Also

[plot.asp](#), [summary.asp](#), [asp](#) (package AdaptFit)

Examples

```
library(SemiPar)
data(fossil)
attach(fossil)
fit <- asp2(strontium.ratio~f(age, basis="tps"))
newdata.age <- data.frame(age=c(90,100,110,120,130))
preds <- predict(fit,newdata=newdata.age,se=TRUE)
print(preds)

# Use predict to avoid centering of smooths in case of scatterplot
# smoothing
fit <- asp2(strontium.ratio~f(age,basis="tps"))
newdata.age <- data.frame(age=seq(90,130,length.out=50) )
preds <- predict(fit,newdata=newdata.age,se=TRUE)
plot(age,strontium.ratio)
lines(newdata.age$age,preds$fit,col="red")
lines(unlist(newdata.age),preds$fit+2*preds$se,col="blue")
lines(unlist(newdata.age),preds$fit-2*preds$se,col="green")
```

residuals.asp

Residuals for semiparametric regression.

Description

Extracts residuals from a semiparametric regression fit object.

Usage

```
## S3 method for class 'asp'
residuals(object,...)
```

Arguments

object a fitted asp object as produced by `asp2()`.
 ... other possible arguments.

Details

Extracts residuals from a semiparametric regression fit object. The residuals are defined to be the difference between the response variable and the fitted values.

Value

The vector of residuals.

See Also

[plot.asp](#), [predict.asp](#), [summary.asp](#), [fitted.asp](#)
[asp](#) (package `AdaptFit`)

Examples

```
library(SemiPar)
data(fossil)
attach(fossil)
fit <- asp2(strontium.ratio~f(age))
plot(age,residuals(fit))
abline(0,0)
```

 scbM

Calculate simultaneous confidence bands for penalized splines

Description

Calculates simultaneous (uniform) confidence bands for the mixed model representation of penalized splines based on volume-of-tube formula.

Usage

```
scbM(object,select=NULL,drv=0,level=0.95,div=1000,
      calc.stdev=TRUE,bayes=FALSE)
```

Arguments

object an asp object.
 select vector specifying which curves in an additive model should be considered. If NULL, all curves are considered.
 drv the derivative order. Defaults to 0, i.e. the estimated function itself is plotted. First and second derivatives are supported.

level	level of confidence.
div	precision for the integral used for calculation of the length of the curve, default is 1000.
calc.stdev	TRUE to compute standard deviation and confidence bands for each value of the covariates. Computationally intensive for large data sets. Use plot.scbm() or plot.asp() to compute standard deviation and bounds only for a grid. If FALSE only critical values are computed.
bayes	FALSE for confidence bands with (approximate) frequentist coverage probability, TRUE for (approximate) Bayesian coverage probability. See Krivobokova et al. (2010) for details.

Details

Returns a scbm object and prints critical values. The resulting confidence bands have (approximate) frequentist coverage probabilities with automatic bias correction (see references). Makes use of the libtube library by Catherine Loader (see package [locfit](#)).

Value

A list object of class scbm containing

aspobject	an asp object.
drv	the derivative order.
crit	a list of critical values.
sigma2	the variance of the residuals.
cov.coef	a list of covariance matrices of spline coefficients in the mixed model framework.
Stdev	the standard deviations of estimates. Only given if calc.stdev=TRUE.
fitted	a list of fitted values. Only given if calc.stdev=TRUE.
lcb	a list of lower bounds of confidence bands. Only given if calc.stdev=TRUE.
ucb	a list of upper bounds of confidence bands. Only given if calc.stdev=TRUE.
...	further

Author(s)

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References

- Krivobokova, T., Kneib, T., and Claeskens, G. (2010)
Simultaneous confidence bands for penalized spline estimators. *Journal of the American Statistical Association*, 105(490):852-863.
- Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition. *Journal of the American Statistical Association*, 107(500): 1286-1296.

Examples

```

## Not run:
beta=function(l,m,x)
return(gamma(l+m)*(gamma(l)*gamma(m))^-1)*x^(l-1)*(1-x)^(m-1))
f1 = function(x) return((0.6*beta(30,17,x)+0.4*beta(3,11,x))*1/0.958)
f2 = function(z) return((sin(2*pi*(z-0.5))^2)*1/.3535)
f3 = function(z)
return((exp(-400*(z-0.6)^2)+
5/3*exp(-500*(z-0.75)^2)+2*exp(-500*(z-0.9)^2))*1/0.549)
center=function(x) return(x-mean(x))

set.seed(1)
N <- 500
x1 = runif(N,0,1)
x2 = runif(N,0,1)
x3 = runif(N,0,1)

kn1 <- default.knots(x1,40)
kn2 <- default.knots(x2,40)
kn3 <- default.knots(x3,40)
kn.var3 <- default.knots(kn3,5)

y <- f1(x1)+f2(x2)+f3(x3)+0.3*rnorm(N)

# fit model with last smoothing parameter adaptive
fit2b= asp2(y~f(x1, basis="os", degree=3, knots=kn1, adap=FALSE)
+f(x2, basis="os", degree=3, knots=kn2, adap=FALSE)
+f(x3, basis="os", degree=3, knots=kn3, adap=TRUE,
var.knots=kn.var3, var.basis="os", var.degree=3),
niter = 20, niter.var = 200)

# compute 95
# You could skip this and use "fit2b" instead of "scb2b" later on,
# however, if N is large, computing the SCBs various times can take
# some time if you don't need fitted values and bounds for all covariate points
# (can be computationally intensive due to large matrix dimensions),
# set calc.stdev=F such that these are not computed.
scb2b<- scbM(fit2b,calc.stdev=FALSE)
plot(scb2b,pages=1)

# plot first derivative of f(x1)
scb2bdrv<- scbM(fit2b,drv=1,calc.stdev=FALSE)
plot(scb2bdrv,select=1)
#the following would give the same result
#plot(fit2b,select=1,drv=1)
#different style
plot(scb2bdrv,select=1,scb.lty="blank", shade=TRUE,
shade.col="steelblue")

## End(Not run)

```

sigma	<i>Extract estimated varying residual variance</i>
-------	--

Description

Extracts the estimated varying residual variance on basis of an object created by aspHetero().

Usage

```
sigma(object)
```

Arguments

object an object created by aspHetero().

Author(s)

Manuel Wiesenfarth <m.wiesenfarth at dkfz de>

Examples

```
#see aspHetero()
```

summary.asp	<i>Semiparametric regression summary</i>
-------------	--

Description

Takes a fitted asp object produced by asp2() and summarises the fit. This function is extensively based on the summary function in package SemiPar.

Usage

```
## S3 method for class 'asp'
summary(object, test1=FALSE, test2=FALSE, signif=0.05, ...)
```

Arguments

object	a fitted asp object as produced by asp2().
test1	TRUE in order to include a test for significance of a nonparametrically estimated effect. The test corresponds to checking whether the zero line is entirely inside the simultaneous confidence band.
test2	TRUE in order to include the nonparametric specification test proposed in Wiesenfarth et al. (2012). Only works with B-splines. The function under the null hypothesis is a polynomial of degree q-1 where q is the penalty order.
signif	the significance level.
...	other arguments.

Details

Produces tables for the linear (parametric) and non-linear (nonparametric) components. The linear table provides coefficient estimates, standard errors and p-values. The non-linear table provides degrees of freedom values and other information.

Value

The function generates summary tables.

References

Ruppert, D., Wand, M.P. and Carroll, R.J. (2003)
Semiparametric Regression Cambridge University Press.
<http://stat.tamu.edu/~carroll/semiregbook/>

Ganguli, B. and Wand, M.P. (2005)
SemiPar 1.0 Users' Manual.
<http://www.uow.edu.au/~mwand/SPmanu.pdf>

Wiesenfarth, M., Krivobokova, T., Klasen, S., Sperlich, S. (2012).
Direct Simultaneous Inference in Additive Models and its Application to Model Undernutrition.
Journal of the American Statistical Association, 107(500): 1286-1296.

See Also

[plot.asp](#), [predict.asp](#)

[asp](#) (package AdaptFit)

Examples

```
## Not run:  
library(SemiPar)  
data(onions)  
attach(onions)  
log.yield <- log(yield)  
fit <- asp2(log.yield~location+f(dens, degree=c(3,2)))  
summary(fit, test1=TRUE, test2=TRUE)  
  
## End(Not run)
```

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