

Package ‘switchnpreg’

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Type Package

Title Switching nonparametric regression models for a single curve and functional data

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Description Functions for estimating the parameters from the latent state process and the functions corresponding to the J states as proposed by De Souza and Heckman (2013).

License GPL-3

Depends MASS, splines, fda

Imports expm, HiddenMarkov

NeedsCompilation no

Repository CRAN

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switchnpreg *Fit a switching nonparametric regression model*

Description

Estimate the parameters of a switching nonparametric regression model using the EM algorithm as proposed by De Souza and Heckman (2013). The package allows two different estimation approaches (Bayesian and penalized log-likelihood) and two different types of hidden states (iid and Markov). The smoothing parameters are chosen by cross-validation. Standard errors for the estimates of the parameters governing the distribution of the state process are also provided.

Usage

```
switchnpreg(x, y, f, alpha, sigma2, lambda, ...,
            method = c("pl", "bayes"), var.equal = TRUE,
            z.indep = TRUE, eps.cv, eps.em, maxit.cv, maxit.em)
```

Arguments

<code>x</code>	The sequence of covariates x_1, \dots, x_n .
<code>y</code>	The sequence of response variables y_1, \dots, y_n .
<code>f</code>	The $n \times J$ matrix of initial values for the functions, where column j corresponds to the function f_j .
<code>alpha</code>	The initial values for the parameters of the latent state process. If the latent states are iid alpha is a vector containing the initial mixing proportions p_j for $j = 1, \dots, J$. If the latent states follow a Markov structure then alpha is a list of two components: A and PI, where A is the initial $J \times J$ matrix of transition probabilities A and PI is the initial J -vector of initial probabilities.
<code>sigma2</code>	The initial J -vector of regression error variances.
<code>lambda</code>	The initial J -vector of smoothing parameters.
<code>...</code>	Optional arguments to parameter update functions.
<code>method</code>	Character string 'pl' or 'bayes' to choose whether the model is fitted using the penalized log-likelihood approach or the Bayesian approach, respectively.
<code>var.equal</code>	Logical indicating whether σ_j^2 are equal for all j .
<code>z.indep</code>	Logical indicating whether the hidden states z_i, \dots, z_n are considered iid or Markovian.
<code>eps.cv</code>	Convergence value for the cross-validation procedure.
<code>eps.em</code>	Convergence value for the EM algorithm.
<code>maxit.cv</code>	Maximum number of iterations of the EM+CV procedure.
<code>maxit.em</code>	Maximum number of iterations of each EM loop.

Value

A list with following elements:

<code>current</code>	The final estimate of θ , represented as a list with the elements named after the respective model parameter: f The final function estimates. sigma2 The final variance estimates. alpha The final estimates for the parameters of the latent state process. pij The matrix of size $n \times J$ with ij -th element giving the final estimate of $p(z_i = j y, \theta)$.
<code>lambda</code>	Chosen smoothing parameters.
<code>iter.cv</code>	Number of iterations of the EM+CV procedure.
<code>stderr</code>	Standard errors for the parameter estimates of the latent state process.

Author(s)

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References

de Souza and Heckman (2013), “Switching nonparametric regression models and the motorcycle data revisited”, submitted for peer review. Available at [arXiv.org](https://arxiv.org/abs/1305.2227v2), article-id: arXiv:1305.2227v2.

See Also

demo(simulated_data_indep_example), demo(simulated_data_Markov_example)

Examples

```
## The motorcycle data set revisited ##

x <- MASS::mcycle$times
set.seed(30)
x[duplicated(x)] <- round(jitter(x[duplicated(x)]),3)

y <- MASS::mcycle$accel

n <- length(y)

spline_fit <- smooth.spline(x, y)

## set up the initial functions
f.initial <- t(apply(as.matrix(spline_fit$y), 1,
                    `+`, c(30, 0, -30)))
J <- ncol(f.initial)
sig2 <- rep((sum((y-predict(spline_fit, x)$y)^2) / (n - spline_fit$df))/J, J)

## B and R parameters for penalized log-likelihood method
basis <- create.bspline.basis(range(x), nbasis = 40)
B <- getbasismatrix(x, basis)
R <- getbasispenalty(basis)

estimates <- switchnpreg(x = x, y = y,
                        f = f.initial,
                        alpha = rep(1, J) / J,
                        sigma2 = sig2,
                        lambda = rep(.5, J),
                        B = B, R = R,
                        var.equal = FALSE,
                        interval = log(c(1E-4, 1E3)),

                        eps.cv = rep(1E-1, J),
                        eps.em = rep(c(1E-1, 1E-2, 1E-3), each = J),
                        maxit.cv = 10,
                        maxit.em = 100)
```

```
plot(x, y, ylim = c(-150,90),
      ylab = 'Head acceleration',
      xlab = 'Time')
matlines(x, estimates$current$f, type='l', lty = 1, col = 1:J)
matlines(sort(x), f.initial, lty = 2, col = 'gray')
```

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