

# Package ‘foser’

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**Title**

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**Description**

**Imports** Rcpp (>= 0.12.14), gibber, dsp, truncdist, MASS

**License** What license is it under?

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 6.0.1

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fosr	<i>MCMC Sampling Algorithm for the Function-on-Scalars Regression Model</i>
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**Description**

Runs the MCMC for the function-on-scalars regression model based on an FDLM-type expansion. Here we assume the factor regression has independent errors, which allows for subject-specific random effects, as well as some additional default conditions.

**Usage**

```
fosr(Y, tau, X = NULL, K = NULL, nsave = 1000, nburn = 1000,
     nskip = 3, mcmc_params = list("beta", "fk", "alpha", "sigma_e",
                                   "sigma_g"), computeDIC = TRUE)
```

**Arguments**

Y	the $n \times m$ data observation matrix, where $n$ is the number of subjects and $m$ is the number of observation points (NAs allowed)
tau	the $m \times d$ matrix of coordinates of observation points
X	the $n \times p$ matrix of predictors; if NULL, only include an intercept
K	the number of factors; if NULL, use SVD-based proportion of variability explained
nsave	number of MCMC iterations to record
nburn	number of MCMC iterations to discard (burn-in)
nskip	number of MCMC iterations to skip between saving iterations, i.e., save every (nskip + 1)th draw
mcmc_params	named list of parameters for which we store the MCMC output; must be one or more of <ul style="list-style-type: none"> <li>• "beta" (factors)</li> <li>• "fk" (loading curves)</li> <li>• "alpha" (regression coefficients)</li> <li>• "mu_k" (intercept term for factor <math>k</math>)</li> <li>• "sigma_e" (observation error SD)</li> <li>• "sigma_g" (random effects SD)</li> <li>• "Yhat" (fitted values)</li> </ul>
computeDIC	logical; if TRUE, compute the deviance information criterion DIC and the effective number of parameters $p_d$

**Value**

A named list of the `nsave` MCMC samples for the parameters named in `mcmc_params`

**Note**

If  $nm$  is large, then storing all posterior samples for  $\hat{Y}$ , which is  $n_{\text{save}} \times n \times M$ , may be inefficient

**Examples**

```
# Simulate some data:
sim_data = simulate_foser(n = 100, m = 20, p_0 = 100, p_1 = 5)

# Data:
Y = sim_data$Y; X = sim_data$X; tau = sim_data$tau

# Dimensions:
n = nrow(Y); m = ncol(Y); p = ncol(X)

# Run the FOSR:
out = foser(Y = Y, tau = tau, X = X, K = 6, mcmc_params = list("fk", "alpha", "Yhat"))

# Plot a posterior summary of a regression function, say j = 3:
j = 3; post_alpha_tilde_j = get_post_alpha_tilde(out$fk, out$alpha[,j])
plot_curve(post_alpha_tilde_j, tau = tau)
# Add the true curve:
lines(tau, sim_data$alpha_tilde_true[,j], lwd=6, col='green', lty=6)

# Plot the loading curves:
plot_flc(out$fk, tau = tau)

# Plot the fitted values for a random subject:
i = sample(1:n, 1)
plot_fitted(y = Y[i,], mu = colMeans(out$Yhat[,i,]),
            postY = out$Yhat[,i,], y_true = sim_data$Y_true[i,], t01 = tau)
```

foser\_gbpv

*Compute Global Bayesian P-Values***Description**

Given posterior samples for the loading curves  $fk$  and the regression coefficient factors  $\alpha$ , compute Global Bayesian P-Values for all regression coefficient functions

**Usage**

```
foser_gbpv(post_fk, post_alpha)
```

**Arguments**

<code>post_fk</code>	$N_{\text{sims}} \times m \times K$ matrix of posterior draws of the loading curve matrix
<code>post_alpha_j</code>	$N_{\text{sims}} \times p \times K$ matrix of posterior draws of the regression coefficient factors

**Value**

$p \times 1$  vector of Global Bayesian P-Values

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foser_select	<i>Decoupling shrinkage and selection for function-on-scalars regression</i>
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**Description**

For a functional response and scalar predictors, construct a posterior summary that balances predictive accuracy and sparsity. Given posterior draws of regression coefficients (or coefficient functions) from a FOSR model, use a suitably-defined loss function to select important variables for prediction.

**Usage**

```
foser_select(X, post_alpha, post_trace_sigma_2, weighted = TRUE,
             alpha_level = 0.1, remove_int = TRUE, include_plot = TRUE,
             include_model_list = FALSE)
```

**Arguments**

X	$n \times p$ matrix of predictors
post_alpha	$N_{\text{sims}} \times p \times K$ array of $N_{\text{sims}}$ posterior draws of the $p$ predictors for each of $K$ factors
post_trace_sigma_2	$N_{\text{sims}} \times 1$ vector of posterior draws of the trace of the (marginal) covariance (see below for details)
weighted	logical; if TRUE, use weighted group lasso (recommended)
alpha_level	coverage for the credible interval on the proportion of variance explained
remove_int	logical; if TRUE, remove the intercept term from model comparisons
include_plot	logical; if TRUE, include a plot showing proportion of variability explained against model size
include_model_list;	if TRUE, include model_list in return—a boolean matrix of models of different sizes suggested by DSS

**Value**

alpha\_dss a  $p \times K$  matrix of (sparse) regression coefficients; if include\_model\_list is TRUE, return a list of alpha\_dss and model\_list, a boolean matrix of possible different sized models

**Note**

This function is value for the regression functions (m-dimensional) as well as the regression factors (K-dimensional). Since  $K \ll m$ , the latter is much faster.

The matrix of predictors,  $X$ , may be different from the given matrix in the data; i.e., we may have a different set of design points for prediction.

post\_trace\_sigma\_2 is the (posterior samples of) the trace of the error covariance matrix jointly across subjects  $i=1, \dots, n$  and observations  $j=1, \dots, m$ , after marginalizing out the random effects  $\gamma_{ik}$ . This is given by  $nm \times \sigma_e^2 + \sum_{ik} \sigma_{\gamma_{ik}}^2$ , where the second term is necessary only when random effects are included in the model AND integrated over in the predictive distribution.

**Examples**

```
# Simulate some data:
sim_data = simulate_fosr(n = 100, m = 20, p_0 = 100, p_1 = 5)

# Data:
Y = sim_data$Y; X = sim_data$X; tau = sim_data$tau

# Dimensions:
n = nrow(Y); m = ncol(Y); p = ncol(X)

# Run the FOSR:
out = fosr(Y = Y, tau = tau, X = X, K = 6, mcmc_params = list("fk", "alpha", "Yhat"))

# Run the DSS:
alpha_dss = fosr_select(X = X,
                        post_alpha = out$alpha,
                        post_trace_sigma_2 = n*m*out$sigma_e^2 + apply(out$sigma_g^2, 1, sum))
# Variables selected:
(select_dss = which(apply(alpha_dss, 1, function(x) any(x != 0))))
```

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get_post_alpha_tilde	<i>Compute the posterior distribution for the regression coefficient functions</i>
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**Description**

Given posterior samples for the loading curves  $fk$  and the regression coefficient factors  $\alpha_j$  for a predictor  $j$ , compute the posterior distribution of the corresponding regression coefficient function.

**Usage**

```
get_post_alpha_tilde(post_fk, post_alpha_j)
```

**Arguments**

post\_fk                Nsims x m x K matrix of posterior draws of the loading curve matrix  
 post\_alpha\_j        Nsims x K matrix of posterior draws of the regression coefficient factors

**Value**

Nsims x m matrix of posterior draws of the regression coefficient function

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plot_curve	<i>Plot a curve given posterior samples</i>
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**Description**

Plot the posterior mean, simultaneous and pointwise 95% credible bands for a curve given draws from the posterior distribution

**Usage**

```
plot_curve(post_f, tau = NULL, alpha = 0.05, include_joint = TRUE,
  main = "Posterior Mean and Credible Bands")
```

**Arguments**

post\_f                Ns x m matrix of Ns posterior simulations of the curve at m points  
 tau                   m x 1 vector of observation points  
 alpha                confidence level for the bands  
 include\_joint        logical; if TRUE, include joint bands (as well as pointwise)  
 main                  text for title plot

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plot_factors	<i>Plot the factors</i>
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**Description**

Plot posterior mean of the factors together with the simultaneous and pointwise 95% credible bands.

**Usage**

```
plot_factors(post_beta, subj = NULL)
```

**Arguments**

post\_beta            the Nsims x n x K array of Nsims draws from the posterior distribution of the  
                          n x K matrix of factors, beta  
 subj                  n x 1 vector of subject IDs or labels

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plot_fitted	<i>Plot the Bayesian curve fitted values</i>
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**Description**

Plot the curve posterior means with posterior credible intervals (pointwise and joint), the observed data, and true curves (if known)

**Usage**

```
plot_fitted(y, mu, postY, y_true = NULL, t01 = NULL,
            include_joint_bands = FALSE)
```

**Arguments**

y	the $n \times 1$ vector of time series observations
mu	the $n \times 1$ vector of fitted values, i.e., posterior expectation of the mean
postY	the $nsims \times n$ matrix of posterior draws from which to compute intervals
y_true	the $n \times 1$ vector of points along the true curve
t01	the observation points; if NULL, assume $n$ equally spaced points from 0 to 1
include_joint_bands	logical; if TRUE, compute simultaneous credible bands

**Examples**

```
# FIXME
```

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plot_flg	<i>Plot the factor loading curves</i>
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**Description**

Plot posterior mean of the factor loading curves together with the simultaneous and pointwise 95% credible bands.

**Usage**

```
plot_flg(post_fk, tau = NULL)
```

**Arguments**

post_fk	the $Nsims \times m \times K$ array of $Nsims$ draws from the posterior distribution of the $m \times K$ matrix of FLCs, fk
tau	$m \times 1$ vector of observation points

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simulate_fosr	<i>Simulate a function-on-scalar regression model</i>
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### Description

Simulate data from a function-on-scalar regression model, allowing for subject-specific random effects. The predictors are multivariate normal with mean zero and covariance  $\text{corr}^{\text{abs}(j1-j2)}$  for correlation parameter  $\text{corr}$  between predictors  $j1$  and  $j2$ . More predictors than observations ( $p > n$ ) is allowed.

### Usage

```
simulate_fosr(n = 100, m = 50, RSNR = 5, K_true = 4, p_0 = 1000,
  p_1 = 5, sparse_factors = TRUE, corr = 0, perc_missing = 0)
```

### Arguments

<code>n</code>	number of observed curves (i.e., number of subjects)
<code>m</code>	total number of observation points (i.e., points along the curve)
<code>RSNR</code>	root signal-to-noise ratio
<code>K_true</code>	rank of the model (i.e., number of basis functions used for the functional data simulations)
<code>p_0</code>	number of true zero regression coefficients
<code>p_1</code>	number of true nonzero regression coefficients
<code>sparse_factors</code>	logical; if TRUE, then for each nonzero predictor $j$ , sample a subset of $k=1:K\_true$ factors to be nonzero#'
<code>corr</code>	correlation parameter for predictors
<code>perc_missing</code>	percentage of missing data (between 0 and 1); default is zero

### Value

a list containing the following:

- `Y`: the simulated  $n \times m$  functional data matrix
- `X`: the simulated  $n \times p$  design matrix
- `tau`: the  $m$ -dimensional vector of observation points
- `Y_true`: the true  $n \times m$  functional data matrix (w/o noise)
- `alpha_tilde_true` the true  $m \times p$  matrix of regression coefficient functions
- `alpha_arr_true` the true  $K\_true \times p$  matrix of regression coefficient factors
- `Beta_true` the true  $n \times K\_true$  matrix of factors
- `F_true` the true  $m \times K\_true$  matrix of basis (loading curve) functions
- `sigma_true` the true observation error standard deviation



**Note**

The basis functions (or loading curves) are orthonormalized polynomials, so large values of `K_true` are not recommended.

**Examples**

```
# Example: simulate FOSR
sim_data = simulate_fosr(n = 100, m = 20, p_0 = 100, p_1 = 5)
Y = sim_data$Y; X = sim_data$X; tau = sim_data$tau
```

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