

# Package ‘midasml’

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**Type** Package

**Title** Estimation and Prediction Methods for High-Dimensional Mixed Frequency Time Series Data

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**Description** The ‘midasml’ estimation and prediction methods for high dimensional time series regression models under mixed data sampling data structures using structured-sparsity penalties and orthogonal polynomials. For more information on the ‘midasml’ approach see Babii, Ghysels, and Striaukas (2020) <arXiv:2005.14057>. Functions that compute MIDAS data structures were inspired by MIDAS ‘Matlab’ toolbox (v2.3) written by Eric Ghysels.

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midasml-package*Estimation and Prediction Methods for High-Dimensional Mixed Frequency Time Series Data*

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**Description**

The 'midasml' estimation and prediction methods for high dimensional time series regression models under mixed data sampling data structures using structured-sparsity penalties and orthogonal polynomials. For more information on the 'midasml' approach see Babii, Ghysels, and Striaukas (2020) <arXiv:2005.14057>. Functions that compute MIDAS data structures were inspired by MIDAS 'Matlab' toolbox (v2.3) written by Eric Ghysels.

**Details**

midasml package contains functions to run classical MIDAS regression models under several loss functions as well as MIDAS machine learning methods. The main functions to run predictive midasml methods are `midasml_forecast` and `qtarget.sort_midasml`, while to estimate the midasml regressions the main functions are `reg_sgl` and `panel_sgl`.

**Author(s)**

Jonas Striaukas [aut, cre]

Maintainer: Jonas Striaukas <jonas.striaukas@gmail.com>

---

apply\_transform

*Time series matrix transformation*

---

**Description**

Transform time series a matrix based on transformation code.

**Usage**

```
apply_transform(data.in, tcode.all)
```

**Arguments**

data.in      time series matrix.

tcode.all      transformation code vector. 1 - not transformed, 2 - first difference, 3 - second difference, 4 - natural log, 5 - first difference of natural log, 6 - second difference of natural log, 7 - first difference of percent change.

**Value**

transformed time series matrix.

**Examples**

```
data.in <- matrix(rnorm(100*2,1),nrow = 100, ncol = 2)
apply_transform(data.in, tcode = rep(1,times=2))
```

---

beta\_w

*Beta density polynomial weights*

---

**Description**

For a given set of parameters  $\theta$  and the number of high-frequency lags, returns weights implied by Beta density functional form.

**Usage**

```
beta_w(param, dayLag)
```

**Arguments**

- `param` two-dimensional parameter vector  $\theta$ .  
`dayLag` number of high-frequency lags.

**Value**

(normalized) weights vector.

**Author(s)**

Jonas Striaukas

**Examples**

```
param <- c(2,2)
dayLag <- 22
beta_w(param, dayLag)
```

`dateMatch`

*Match dates*

**Description**

Change the date to the beginning of the month date.

**Usage**

```
dateMatch(x, y)
```

**Arguments**

- `x` date vector to match with `y` date vector.  
`y` date vector.

**Value**

changed date vector.

**Author(s)**

Jonas Striaukas

**Examples**

```
x <- seq(as.Date("2020-01-01"), as.Date("2020-12-01"), by = "day")
set.seed(100)
x <- x[-sample(1:336, 100)]
y <- seq(as.Date("2020-01-01"), as.Date("2020-12-01"), by = "month")
dateMatch(x,y)
```

expalmon\_w

*Exponential Almon polynomial weights***Description**

For a given set of parameters  $\theta$  and the number of high-frequency lags, returns weights implied by exponential Almon functional form.

**Usage**

```
expalmon_w(param, dayLag)
```

**Arguments**

- |        |   |
|--------|---|
| param  | two-dimensional parameter vector $\theta$ . |
| dayLag | number of high-frequency lags.              |

**Value**

(normalized) weights vector.

**Author(s)**

Jonas Striaukas

**Examples**

```
param <- c(0.02, -0.2)
dayLag <- 22
expalmon_w(param, dayLag)
```

gb

*Gegenbauer polynomials shifted to [a,b]***Description**

For a given set of points in X, computes the orthonormal Gegenbauer polynomials basis of L2 [a,b] for a given degree and  $\alpha$  parameter. The Gegenbauer polynomials are a special case of more general Jacobi polynomials. In turn, you may get Legendre polynomials from Gegenbauer by setting  $\alpha = 0$ , or Chebychev's polynomials by setting  $\alpha = 1/2$  or  $-1/2$ .

**Usage**

```
gb(degree, alpha, a = 0, b = 1, jmax = NULL, X = NULL)
```

**Arguments**

degree	polynomial degree.
alpha	Gegenbauer polynomials parameter.
a	lower shift value (default - 0).
b	upper shift value (default - 1).
jmax	number of high-frequency lags.
X	optional evaluation grid vector.

**Value**

Psi weight matrix with Gegenbauer functions upto degree.

**Author(s)**

Jonas Striaukas

**Examples**

```
degree <- 3
alpha <- 1
jmax <- 66
gb(degree = degree, alpha = alpha, a = 0, b = 1, jmax = jmax)
```

lb

*Legendre polynomials shifted to [a,b]*

**Description**

For a given set of points in X, computes the orthonormal Legendre polynomials basis of L2 [a,b] for a given degree.

**Usage**

```
lb(degree, a = 0, b = 1, jmax = NULL, X = NULL)
```

**Arguments**

degree	polynomial degree.
a	lower shift value (default - 0).
b	upper shift value (default - 1).
jmax	number of high-frequency lags.
X	optional evaluation grid vector.

**Value**

Psi weight matrix with Legendre functions upto degree.

**Author(s)**

Jonas Striaukas

**Examples**

```
degree <- 3
jmax <- 66
lb(degree = degree, a = 0, b = 1, jmax = jmax)
```

---

macro\_midasml

*GDP nowcasting using midasML approach example data*

---

**Description**

US real GDP, FRED-MD monthly dataset, SPF survey and textual analysis data.

**Usage**

```
data(market_ret)
```

**Format**

A [list](#) object

**Source**

macro\_midasml\$rgdp - [FRED](#)  
macro\_midasml\$md.data - [St. Louis Fed M. W. McCracken website](#)  
macro\_midasml\$text.data - [The Structure of Economic News website](#)  
macro\_midasml\$survey.data - [Philadelphia Fed SPF data](#)

**Examples**

```
data(macro_midasml)
macro_midasml$rgdp.data # GDP data
macro_midasml$md.data # FRED-MD data
macro_midasml$text.data # textual analysis data
macro_midasml$survey.data # SPF data
```

---

market_ret	<i>SNP500 returns</i>
------------	-----------------------

---

**Description**

SNP500 returns

**Usage**

```
data(market_ret)
```

**Format**

A `data.frame` object.a

**Source**

rgdp - FRED

**Examples**

```
data(market_ret)
market_ret$snp500ret
```

---

midasml_forecast	<i>MIDAS ML regression prediction function</i>
------------------	--

---

**Description**

Predicts from a high-dimensional MIDAS model.

**Usage**

```
midasml_forecast(y_in, y_out, x_in, x_out, group_index,
                  gamma_w, y_out_dates, scheme, verbose = FALSE, ...)
```

**Arguments**

y_in	response variable (in-sample).
y_out	response variable (out-of-sample).
x_in	predictor variables (in-sample).
x_out	predictor variables (out-of-sample).
group_index	group membership of each covariate.

gamma_w	sg-LASSO mixing parameter. gamma_w = 1 is LASSO and gamma_w = 0 group LASSO.
y_out_dates	out-of-sample dates.
scheme	prediction scheme. Choices are: expand - expanding window scheme, rolling - rolling window scheme.
verbose	flag to print information.
...	optional parameters to feed into reg_sgl.

## Details

Based on desired computation of the tuning parameter  $\lambda$  (ic/cv), the MIDAS ML regression is fit and predictions are computed based on the chosen scheme (rolling/expanding). The regression function that is fit is

$$\|y - \alpha\iota - X\beta\|_T^2 + 2\lambda\Omega_\gamma(\beta)$$

,

where  $X$  ( $x_{in}$  and  $x_{out}$ ) contains autoregressive lags and MIDAS covariates of sort\_midasml class. The group index vector should be used from data sorting function qtarget.sort\_midasml. The penalty function  $\Omega_\gamma(\cdot)$  is applied on  $\beta$  coefficients and is

$$\Omega_\gamma(\beta) = \gamma|\beta|_1 + (1 - \gamma)\|\beta\|_{2,1},$$

a convex combination of LASSO and group LASSO penalty functions.

## Value

out-of-sample predictions.

## Author(s)

Jonas Striaukas

## Examples

```
data(macro_midasml)
est.start <- as.Date("1990-12-01")
est.end <- as.Date("2017-03-01")
rgdp.data <- macro_midasml$rgdp.data
rgdp.data <- rgdp.data[rgdp.data$DATE<=as.Date("2017-06-01"),]
data <- qtarget.sort_midasml(y.data = rgdp.data, x.macro.data = macro_midasml$md.data,
                            x.real.time = macro_midasml$text.data, x.quarterly_group = macro_midasml$survey.data,
                            x.lag = 12, legendre_degree = 3,
                            horizon = 1, macro_delay = 1, est.start, est.end,
                            standardize = TRUE, group_ar_lags = FALSE, disp.flag = FALSE)
midasml_forecast(y_in = data$y_in, y_out = data$y_out,
                  x_in = data$x_str, x_out = data$x_str_out,
                  group_index = data$group_index, gamma_w = 0.65,
```

---

```
y_out_dates = data$y_out_dates, scheme = "expand",
method_choice = "ic", num_cores = 2)
```

---

**midas\_ardl***ARDL-MIDAS regression***Description**

Estimates and predicts using a single variate ARDL-MIDAS model.

**Usage**

```
midas_ardl(data.y, data.ydate, data.x, data.xdate, x.lag,
y.lag, est.start, est.end, horizon = 1,
polynomial = c("legendre_w", "beta_w", "rbeta_w", "expalmon_w", "umidas_w", "step_fun"),
scheme = c("fixed", "rolling", "expand"), loss = c("mse", "rq", "als"), ...)
```

**Arguments**

<code>data.y</code>	response variable series.
<code>data.ydate</code>	response variable dates.
<code>data.x</code>	predictor variable series.
<code>data.xdate</code>	predictor variable dates.
<code>x.lag</code>	number of high-frequency data lags.
<code>y.lag</code>	number of low-frequency data lags.
<code>est.start</code>	start date of the estimation sample (referenced with <code>data.xdate</code> ).
<code>est.end</code>	end date of the estimation sample (referenced with <code>data.xdate</code> ).
<code>horizon</code>	forecast horizon measured in predictor variable sampling frequency (default set 1 unit ahead).
<code>polynomial</code>	MIDAS lag polynomial specification. Options are: Legendre ( <code>legendre_w</code> ), Beta density ( <code>beta_w</code> ), restricted Beta density ( <code>rbeta_w</code> ), exponential Almon ( <code>expalmon_w</code> ), unrestricted MIDAS ( <code>umidas_w</code> ), step functions ( <code>step_fun</code> ).
<code>scheme</code>	forecasting scheme. Options are: fixed scheme ( <code>fixed</code> ), rolling window scheme ( <code>rolling</code> ), expanding window scheme ( <code>expand</code> ).
<code>loss</code>	loss function. Options are: mean squared error ( <code>mse</code> ), quantile ( <code>rq</code> ), asymmetric least squares ( <code>als</code> ).
<code>...</code>	optional parameters to feed into other functions. <code>forecast.flag</code> - TRUE/FALSE to compute out-of-sample predictions (default TRUE), <code>disp.flag</code> - TRUE/FALSE to display MIDAS data structures (default FALSE) <code>num.evals</code> - number of objective function evaluations using random starting parameter values in the case of non-linear MIDAS polynomial (default 1e4) <code>num.coef</code> - number of best coefficients to use as starting values in nonlinear optimization (default 10)

seed - value used in set.seed for randomly drawing initial starting values around OLS optimal solution  
 profiling - TRUE/FALSE to use MIDAS parameter profiling, coded only for rbeta\_w polynomial, (default FALSE)  
 step\_idx - index of step function lags. If step\_fun is used as a polynomial, it is best to specify this option too, otherwise, the program figures out the sampling frequency ratio and computes step\_idx accordingly (message is displayed in this case)  
 legendre\_degree - degree of Legendre polynomials. If legendre\_w is used as a polynomial, it is best to specify this option too, otherwise, the value is set to 3 (message is displayed in this case)  
 tau - quantile level for als and rq regressions. If either als or rq loss is used, this option must be specified, program stops if no value is provided.

## Details

Several polynomial functional forms are available (input variable polynomial):

- beta\_w: Beta polynomial
- rbeta\_w: restricted Beta polynomial
- expalmon\_w: Exp Almon polynomial
- umidas\_w: unrestricted lags (U-MIDAS)
- step\_fun: polynomial with step functions
- legendre\_w: Legendre polynomials

different forecasting schemes (input variable scheme):

- fixed: fixed scheme
- rolling: rolling window
- expand: expanding window

and different loss functions (input variable loss)

- mse: least squares
- als: asymmetric least squares
- rq: quantile.

The ARDL-MIDAS model is:

$$y_t = \mu + \sum_p \rho_p y_{t-p} + \beta \sum_j \omega_j(\theta) x_{t-1}$$

where  $\mu, \beta, \theta, \rho_p$  are model parameters, p is number of low-frequency and  $\omega$  is the weight function.

## Value

returns midas\_ardl list which contains parameter estimates, in- and out-of-sample statistics and predictions, and some information about the specification of the method used.

## Author(s)

Jonas Striaukas

## Examples

```
data(us_rgdp)
rgdp <- us_rgdp$rgdp
cfnai <- us_rgdp$cfnai
rgdp[-1, 2] <- ((rgdp[-1, 2]/rgdp[-dim(rgdp)[1], 2])^4-1)*100
rgdp <- rgdp[-1, ]
data.y <- rgdp[,2]
data.ydate <- rgdp[,1]
est.start <- as.Date("1990-01-01")
est.end <- as.Date("2002-03-01")
data.x <- cfnai[,2]
data.xdate <- cfnai[,1]
midas_ardl(data.y, data.ydate, data.x, data.xdate,
            x.lag = 12, y.lag = 4, est.start, est.end, horizon = 1,
            polynomial = "legendre_w", legendre_degree = 3)
```

midas\_dl

*DL-MIDAS regression*

## Description

Estimates and predicts using a single variate DL-MIDAS model.

## Usage

```
midas_dl(data.x, data.xdate, data.y, data.ydate, x.lag, est.start, est.end, horizon = 1,
          polynomial = c("legendre_w", "beta_w", "rbeta_w", "expalmon_w", "umidas_w", "step_fun"),
          scheme = c("fixed", "rolling", "expand"), loss = c("mse", "rq", "als"), ...)
```

## Arguments

data.x	predictor variable series.
data.xdate	predictor variable dates.
data.y	dependent variable series (can leave unspecified, see <code>midas_gen</code> option).
data.ydate	dependent variable dates (can leave unspecified, see <code>midas_gen</code> option).
x.lag	number of high-frequency data lags.
est.start	start date of the estimation sample (referenced with <code>data.xdate</code> ).
est.end	end date of the estimation sample (referenced with <code>data.xdate</code> ).
horizon	forecast horizon measured in predictor variable sampling frequency (default set 1 unit ahead).
polynomial	MIDAS lag polynomial specification. Options are: Legendre ( <code>legendre_w</code> ), Beta density ( <code>beta_w</code> ), restricted Beta density ( <code>rbeta_w</code> ), exponential Almon ( <code>expalmon_w</code> ), unrestricted MIDAS ( <code>umidas_w</code> ), step functions ( <code>step_fun</code> ).
scheme	forecasting scheme. Options are: fixed scheme ( <code>fixed</code> ), rolling window scheme ( <code>rolling</code> ), expanding window scheme ( <code>expand</code> ).

**loss** loss function. Options are: mean squared error (mse), quantile (rq), asymmetric least squares (als).

**...** optional parameters to feed into other functions:  
**forecast.flag** - TRUE/FALSE to compute out-of-sample predictions (default TRUE)  
**disp.flag** - TRUE/FALSE to display MIDAS data structures (default FALSE)  
**num.evals** - number of objective function evaluations using random starting parameter values in the case of non-linear MIDAS polynomial (default 1e4)  
**num.coef** - number of best coefficients to use as starting values in nonlinear optimization (default 10)  
**seed** - value used in set.seed for randomly drawing initial starting values around OLS optimal solution  
**profiling** - TRUE/FALSE to use MIDAS parameter profiling, coded only for rbeta\_w polynomial, (default FALSE)  
**step\_idx** - index of step function lags. If **step\_fun** is used as a polynomial, it is best to specify this option too, otherwise, the program figures out the sampling frequency ratio and computes **step\_idx** accordingly (message is displayed in this case)  
**legendre\_degree** - a degree of legendre polynomials. If **legendre\_w** is used as a polynomial, it is best to specify this option too, otherwise, the value is set to 3 (message is displayed in this case)  
**tau** - quantile level for als and rq regressions. If either als or rq loss is used, this option must be specified, program stops if no value is provided  
**midas\_gen** - option on how to generate the low-frequency variable. **from\_hf** - computes from high-frequency variable (see **mixed\_freq\_data\_mhorizon**, aggregation method could be specified as an additional input) or **as\_ref** - computes MIDAS data structures using low-frequency variable (default 'from\_hf').

## Details

Several polynomial functional forms are available (input variable **polynomial**):

- **beta\_w**: Beta polynomial
- **rbeta\_w**: restricted Beta polynomial
- **expalmon\_w**: Exp Almon polynomial
- **umidas\_w**: unrestricted lags (U-MIDAS)
- **step\_fun**: polynomial with step functions
- **legendre\_w**: Legendre polynomials

different forecasting schemes (input variable **scheme**):

- **fixed**: fixed scheme
- **rolling**: rolling window
- **expand**: expanding window

and different loss functions (input variable **loss**)

- **mse**: least squares
- **als**: asymmetric least squares
- **rq**: quantile.

The DL-MIDAS model is:

$$y_t = \mu + \beta \sum_j \omega_j(\theta) x_{t-1}$$

where  $\mu$ ,  $\beta$  and  $\theta$  are model parameters and  $\omega$  is the weight function.

### **Value**

returns `midas_dl` list which contains parameter estimates, in- and out-of-sample statistics and predictions, and some information about the specification of the method used.

### **Author(s)**

Jonas Striaukas

### **Examples**

```
data(market_ret)
data.x <- market_ret$snp500ret
data.xdate <- market_ret$DATE
est.start <- as.Date("2005-01-01")
est.end <- as.Date("2008-12-31")
midas_dl(data.x, data.xdate, x.lag = 5,
          est.start = est.start, est.end = est.end,
          horizon = 1, polynomial = "legendre_w", legendre_degree = 3,
          scheme = "fixed", loss = "mse", midas_gen = "from_hf")
```

**mixed\_freq\_data**

*MIDAS data structure*

### **Description**

Creates a MIDAS data structure for a single high-frequency covariate and a single low-frequency dependent variable.

### **Usage**

```
mixed_freq_data(data.y, data.ydate, data.x, data.xdate, x.lag, y.lag,
                horizon, est.start, est.end, disp.flag = TRUE)
```

### **Arguments**

<code>data.y</code>	n by 1 low-frequency time series data vector.
<code>data.ydate</code>	n by 1 low-frequency time series date vector.
<code>data.x</code>	nm by 1 high-frequency time series data vector.
<code>data.xdate</code>	nm by 1 high-frequency time series date vector.
<code>x.lag</code>	number of high-frequency lags to construct in high-frequency time units.

y.lag	number of low-frequency lags to construct in low-frequency time units.
horizon	forecast horizon relative to data.ydate date in high-frequency time units.
est.start	estimation start date, taken as the first ... .
est.end	estimation end date, taken as the last ... . Remaining data after this date is dropped to out-of-sample evaluation data.
disp.flag	display flag to indicate whether or not to display obtained MIDAS data structure in console.

**Value**

a list of MIDAS data structure.

**Author(s)**

Jonas Striaukas

**Examples**

```
data(us_rgdp)
rgdp <- us_rgdp$rgdp
payems <- us_rgdp$payems
payems[-1, 2] <- log(payems[-1, 2]/payems[-dim(payems)[1], 2])*100
payems <- payems[-1, ]
rgdp[-1, 2] <- ((rgdp[-1, 2]/rgdp[-dim(rgdp)[1], 2])^4-1)*100
rgdp <- rgdp[-1, ]
est.start <- as.Date("1990-01-01")
est.end <- as.Date("2002-03-01")
mixed_freq_data(rgdp[,2], as.Date(rgdp[,1]), payems[,2],
  as.Date(payems[,1]), x.lag = 9, y.lag = 4, horizon = 1,
  est.start, est.end, disp.flag = FALSE)
```

**mixed\_freq\_data\_mhorizon**

*MIDAS data structure*

**Description**

Creates a MIDAS data structure for a single high-frequency covariate and a single low-frequency dependent variable computed from high-frequency covariate (e.g. stock returns).

**Usage**

```
mixed_freq_data_mhorizon(data.x, data.xdate, x.lag, est.start, est.end,
  horizon, disp.flag = TRUE, ...)
```

**Arguments**

<code>data.x</code>	nm by 1 high-frequency time series data vector.
<code>data.xdate</code>	nm by 1 high-frequency time series date vector.
<code>x.lag</code>	number of high-frequency lags to construct in high-frequency time units.
<code>est.start</code>	estimation start date, taken as the first ... .
<code>est.end</code>	estimation end date, taken as the last ... . Remaining data after this date is dropped to out-of-sample evaluation data.
<code>horizon</code>	forecast horizon relative to <code>data.ydate</code> date in high-frequency time units.
<code>disp.flag</code>	display flag to indicate whether or not to display obtained MIDAS data structure in console.
<code>...</code>	an optional parameter aggregation specifying the aggregation method of high-frequency data to get low-frequency target (non-overlapping) <code>sum</code> - sum of high-frequency lags <code>sum&amp;abs</code> - sum of high-frequency lags which after aggregation are taken in absolute value <code>sum&amp;sq</code> - sum of high-frequency lags which after aggregation are taken in squares <code>mean</code> - average of high-frequency lags <code>first_val</code> - the most recent lag value of high-frequency lags.

**Value**

a list of midas data structure.

**Author(s)**

Jonas Striaukas

**Examples**

```
data(market_ret)
data.x <- market_ret$snp500ret
data.xdate <- market_ret$DATE
est.start <- as.Date("2005-01-01")
est.end <- as.Date("2017-12-31")
mixed_freq_data_mh horizon(data.x, data.xdate, x.lag = 5, est.start, est.end,
                           horizon = 1, disp.flag = FALSE, aggregation = "sum")
```

`mixed_freq_data_single`

*MIDAS data structure*

**Description**

Creates a MIDAS data structure for a single high-frequency covariate based on low-frequency reference date.

**Usage**

```
mixed_freq_data_single(data.refdate, data.x, data.xdate, x.lag, horizon,
est.start, est.end, disp.flag = TRUE)
```

**Arguments**

data.refdate	n by 1 date vector.
data.x	nm by 1 high-frequency time series data vector.
data.xdate	nm by 1 high-frequency time series date vector.
x.lag	number of high-frequency lags to construct in high-frequency time units.
horizon	forecast horizon relative to data.refdate date in high-frequency time units.
est.start	estimation start date, taken as the first ... .
est.end	estimation end date, taken as the last ... . Remaining data after this date is dropped to out-of-sample evaluation data.
disp.flag	display flag to indicate whether or not to display obtained MIDAS data strcuture in console.

**Value**

a list of midas data structure.

**Author(s)**

Jonas Striaukas

**Examples**

```
data(us_rgdp)
rgdp <- us_rgdp$rgdp
cfnai <- us_rgdp$cfnai
data.refdate <- rgdp$date
data.x <- cfnai$cfnai
data.xdate <- cfnai$date
est.start <- as.Date("1990-01-01")
est.end <- as.Date("2002-03-01")
mixed_freq_data_single(data.refdate, data.x, data.xdate, x.lag = 12, horizon = 1,
est.start, est.end, disp.flag = FALSE)
```

---

monthBegin	<i>Beginning of the month date</i>
------------	------------------------------------

---

**Description**

Change the date to the beginning of the month date.

**Usage**

```
monthBegin(x)
```

**Arguments**

x date value.

**Value**

changed date value.

**Author(s)**

Jonas Striaukas

**Examples**

```
monthBegin(as.Date("2020-05-15"))
```

---

monthEnd	<i>End of the month date</i>
----------	------------------------------

---

**Description**

Change the date to the end of the month date.

**Usage**

```
monthEnd(x)
```

**Arguments**

x date value.

**Value**

changed date value.

**Author(s)**

Jonas Striaukas

**Examples**

```
monthEnd(as.Date("2020-05-15"))
```

panel\_sgl

*Panel sg-LASSO regression model*

**Description**

Fits panel data regression model, random and fixed effects, under sg-LASSO penalty function. Options include random effects and fixed effects models, cross-validation and information criteria for  $\lambda$  penalty parameter selection.

**Usage**

```
panel_sgl(X, Z = NULL, y, index, entity_indices, gamma_w = NULL, l1_factor = NULL,
  l21_factor = NULL, dummies_index = NULL, full_est = NULL,
  regress_choice = c("re", "fe"), method_choice = c("ic", "cv", "initial"),
  nlam = 100, lambdas = NULL, min_frac = NULL, nfolds = 10,
  lambda_choice = c("min", "1se"), ic_choice = c("bic", "aic", "aicc"),
  num_cores = NULL, verbose = FALSE, thresh = NULL,
  outer_thresh = NULL, inner_iter = NULL, outer_iter = NULL)
```

**Arguments**

X	NT by p data matrix, where n, t and p respectively denote the number of individuals, sample size and the number of regressors.
Z	dummies matrix for random effects or fixed effects panel data model. If left unspecified, it is computed based on regress_choice choice.
y	NT by 1 vector of outcome.
index	p by 1 vector indicating group membership of each covariate.
entity_indices	NT by 1 vector of individual indices.
gamma_w	sg-LASSO mixing parameter. $\text{gamma\_w} = 1$ is LASSO and $\text{gamma\_w} = 0$ group LASSO.
l1_factor	$\ell_1$ norm penalty factor for random or fixed effects (default value is zero which means $\alpha$ is left unpenalized in $\ell_1$ norm).
l21_factor	$\ell_1$ norm penalty factor for random or fixed effects (default value is zero which means $\alpha$ is left unpenalized in $\ell_1$ norm).
dummies_index	vector indicating group membership of $\alpha$ (default - no grouping).
full_est	pre-estimated parameters based on full sample and regress_choice for a sequence of $\lambda$ 's.

<code>regress_choice</code>	choose between ‘re’ and ‘fe’. ‘re’ computes random effects regression with sg-LASSO penalty (default). ‘fe’ computes fixed effects regression with sg-LASSO penalty.
<code>method_choice</code>	choose between ‘initial’, ‘ic’ and ‘cv’. ‘initial’ pre-computes initial estimates. ‘ic’ computes solution based on information criteria (BIC, AIC or AICC). ‘cv’ computes solution based on cross-validation (cv).
<code>nlam</code>	number of $\lambda$ ’s to use in the regularization path.
<code>lambdas</code>	user specified sequence of $\lambda$ values for fitting. We recommend leaving this to NULL and letting function to self-select values.
<code>min_frac</code>	the minimum value of the penalty parameter, as a fraction of the maximum value.
<code>nfolds</code>	number of folds of the cv loop.
<code>lambda_choice</code>	choose between ‘min’ and ‘1se’. ‘min’ computes solution that minimizes the cv error. ‘1se’ computes solution such that the cv error is within 1 standard error of the minimum ‘min’.
<code>ic_choice</code>	choose between ‘bic’, ‘aic’ and ‘aicc’. ‘bic’ computes solution that minimizes Bayesian information criterion. ‘aic’ computes solution that minimizes Akaike information criterion. ‘aicc’ computes solution that minimizes Akaike information corrected criterion.
<code>num_cores</code>	number of cores used to compute cv loop.
<code>verbose</code>	flag to print information.
<code>thresh</code>	convergence threshold for change in beta. We recommend leaving this to NULL.
<code>outer_thresh</code>	outer loop convergence threshold. We recommend leaving this to NULL.
<code>inner_iter</code>	the maximum number of inner sg-LASSO loop iterations. We recommend leaving this to NULL.
<code>outer_iter</code>	the maximum number of outer sg-LASSO loop iterations. We recommend leaving this to NULL.

## Details

The sequence of panel data models implied by `lambdas` vector is fit by block coordinate-descent. The objective function is

$$RSS(\alpha, \beta)/NT + 2\lambda * \Omega_\gamma(\beta),$$

where  $RSS(\alpha, \beta)$  is either random or fixed effects model fit. The penalty function  $\Omega_\gamma(\cdot)$  is applied on coefficients of time-varying covariates  $\beta$  and is

$$\Omega_\gamma(\beta) = \gamma|\beta|_1 + (1 - \gamma)\|\beta\|_{2,1},$$

a convex combination of LASSO and group LASSO penalty functions. There are additional options to apply sg-LASSO structures on fixed effects coefficient vector  $\alpha$  (see description of input variables for more details).

## Value

Parameter estimates of panel data regression model under sg-LASSO penalty.

**Author(s)**

Jonas Striaukas

**Examples**

```
# simulate DGP
set.seed(1)
t <- 10; n = 5; p = 20; size.groups = 4
index <- ceiling(1:p / size.groups)
X <- matrix(rnorm(n * t * p), ncol = p, nrow = n*t)
beta <- c(5,4,3,2,1)
y <- X[,1:5] %*% beta + 5*rnorm(n*t)
entity_indices <- sort(rep(1:n,times=t-1))
panel_sgl(X = X, Z = NULL, y = y, index = index,
entity_indices = entity_indices, gamma_w = 1,
regress_choice = "fe", method_choice = "initial",
num_cores = 2, verbose = FALSE)
```

**plot\_weights**

*MIDAS weights plot function*

**Description**

Based on specification in obj, plots a basic R figure of estimated MIDAS weights.

**Usage**

```
plot_weights(obj)
```

**Arguments**

obj	midas_ardl or midas_dl object with parameter estimates and model specification inputs.
-----	--

**Details**

MIDAS regression pecifcation is picked up from obj, see `midas_dl` or `midas_ardl` function descriptions for more details.

**Value**

returns R figure of estimated MIDAS weights.

**Author(s)**

Jonas Striaukas

## Examples

```

data(us_rgdp)
rgdp <- us_rgdp$rgdp
cfnai <- us_rgdp$cfnai
rgdp[-1, 2] <- ((rgdp[-1, 2]/rgdp[-dim(rgdp)[1], 2])^4-1)*100
rgdp <- rgdp[-1, ]
data.y <- rgdp[,2]
data.ydate <- rgdp[,1]
est.start <- as.Date("1990-01-01")
est.end <- as.Date("2002-03-01")
data.x <- cfnai[,2]
data.xdate <- cfnai[,1]
fit <- midas_ardl(data.y, data.ydate, data.x, data.xdate,
                     x.lag = 12, y.lag = 4, est.start, est.end, horizon = 1,
                     polynomial = "legendre_w", legendre_degree = 3)
plot_weights(obj = fit$est.obj)

```

**predict.panel\_sgl**      *Computes prediction for the sg-LASSO panel regression model*

## Description

Computes prediction for the sg-LASSO panel regression model

## Usage

```

## S3 method for class 'panel_sgl'
predict(object, newX, newZ = NULL, regress_choice = c("re", "fe"), ...)

```

## Arguments

object	fit object from panel_sgl.
newX	matrix of out-of-sample covariate observations.
newZ	optional matrix of dummies for panel data model.
regress_choice	choose between ‘re’ and ‘fe’. Must be consistent with object.
...	currently ignored optional parameters.

## Value

a list of these variables:  
 pred - overall prediction.  
 predZ - dummies prediction.  
 predX - covariates prediction.

**Author(s)**

Jonas Striaukas

**Examples**

```
set.seed(1)
t <- 10; n = 5; p = 20; size.groups = 4
index <- ceiling(1:p / size.groups)
X <- matrix(rnorm(n * t * p), ncol = p, nrow = n*t)
beta <- c(5,4,3,2,1)
y <- X[,1:5] %*% beta + 5*rnorm(n*t)
Z <- kronecker(diag(n), rep(1, times = t))
entity_indices <- sort(rep(1:n,times=t-1))
fit <- panel_sgl(X = X, Z = Z, y = y, index = index,
                   entity_indices = entity_indices, gamma_w = 1,
                   regress_choice = "fe", method_choice = "ic",
                   num_cores = 2, verbose = FALSE)
predict.panel_sgl(object = fit, newX = X, newZ = Z, regress_choice = "fe")$pred
```

**predict.reg\_sgl**

*Computes prediction for the sg-LASSO linear regression*

**Description**

Computes prediction for the sg-LASSO linear regression

**Usage**

```
## S3 method for class 'reg_sgl'
predict(object, newX, ...)
```

**Arguments**

object	fit object from sglassofit.
newX	matrix of out-of-sample covariate observations.
...	currently ignored optional parameters.

**Value**

a list of these variables:  
 pred - overall prediction.  
 predZ - intercept prediction.  
 predX - covariates prediction.

**Author(s)**

Jonas Striaukas

**Examples**

```
set.seed(1)
x <- matrix(rnorm(100 * 20), 100, 20)
y <- rnorm(100)
index <- 1:20
fit <- reg_sgl(X = x, y = y, index = index, gamma_w = 1, method_choice = "ic",
                 num_cores = 2, verbose = FALSE)
predict.reg_sgl(object = fit, newX = x)
```

*qtarget.sort\_midasml*    *High-dimensional mixed frequency data sort function*

**Description**

Sorts high-dimensional mixed frequency data for quarterly target variable.

**Usage**

```
qtarget.sort_midasml(y.data, x.macro.data = NULL, x.real.time = NULL,
                      x.quarterly_group = NULL, x.lag = NULL, legendre_degree, horizon,
                      macro_delay = 1, est.start, est.end, standardize = TRUE, group_ar_lags = FALSE,
                      real_time_predictions = FALSE, disp.flag = TRUE)
```

**Arguments**

<i>y.data</i>	response variable data.
<i>x.macro.data</i>	macro data which is not real-time, i.e. is used with publication delay defined in <i>macro_delay</i> .
<i>x.real.time</i>	real-time data.
<i>x.quarterly_group</i>	quarterly data currently taken as real-time data.
<i>x.lag</i>	single value or vector of size of the total number of variables defining the number of lags for each high-frequency variable in <i>x.macro.data</i> , <i>x.real.time</i> .
<i>legendre_degree</i>	single value or vector of size of the total number of variables defining the polynomial degree for each high-frequency variable in <i>x.macro.data</i> , <i>x.real.time</i> .
<i>horizon</i>	forecast horizon relative to <i>y.data</i> date column in high-frequency time units.
<i>macro_delay</i>	number of months that macro series in <i>x.macro.data</i> are delayed.
<i>est.start</i>	estimation start date, taken as the first ... .

est.end	estimation end date, taken as the last ... . Remaining data after this date is dropped to out-of-sample evaluation data.
standardize	TRUE/FALSE to standardize high-frequency covariates in high-frequency units.
group_ar_lags	TRUE/FALSE to group AR lags.
real_time_predictions	TRUE/FALSE in case real-time data is used for predictions
disp.flag	display flag to indicate whether or not to display obtained MIDAS data structure in console.

## Details

Sorts high-dimensional mixed frequency data for quarterly target variable read to be inputed into midasml\_forecast function to produce nowcasts & forecasts.

## Value

MIDAS covariates and group memberships based on desired specification.

## Author(s)

Jonas Striaukas

## Examples

```
data(macro_midasml)
est.start <- as.Date("1990-12-01")
est.end <- as.Date("2017-03-01")
rgdp.data <- macro_midasml$rgdp.data
rgdp.data <- rgdp.data[rgdp.data$DATE<=as.Date("2017-06-01"),]
qtarget.sort_midasml(y.data = rgdp.data, x.macro.data = macro_midasml$md.data,
                      x.real.time = macro_midasml$text.data, x.quarterly_group = macro_midasml$survey.data,
                      x.lag = 12, legendre_degree = 3,
                      horizon = 1, macro_delay = 1, est.start, est.end,
                      standardize = TRUE, group_ar_lags = FALSE, disp.flag = FALSE)
```

## Description

For a given set of parameters  $\theta$  and the number of high-frequency lags, returns weights implied by Restricted Beta density functional form.

## Usage

rbeta\_w(param, dayLag)

**Arguments**

- param one-dimensional parameter vector  $\theta$ .  
 dayLag number of high-frequency lags.

**Value**

(normalized) weights vector.

**Author(s)**

Jonas Striaukas

**Examples**

```
param <- 3
dayLag <- 22
rbeta_w(param, dayLag)
```

reg\_sgl

*Linear sg-LASSO regression*

**Description**

Fits sg-LASSO least squares regression model. Options include cross-validation and information criteria for  $\lambda$  penalty parameter selection.

**Usage**

```
reg_sgl(X, y, index, gamma_w = NULL, full_est = NULL,
method_choice = c("ic", "cv", "initial"), nlam = 100, lambdas = NULL,
min_frac = NULL, nfolds = 10, lambda_choice = c("min", "1se"),
ic_choice = c("bic", "aic", "aicc"), num_cores = NULL, verbose = FALSE,
thresh = NULL, outer_thresh = NULL, inner_iter = NULL, outer_iter = NULL)
```

**Arguments**

- X T by p data matrix, where t and p respectively denote the sample size and the number of regressors.  
 y T by 1 vector of outcome.  
 index p by 1 vector indicating group membership of each covariate.  
 gamma\_w sg-LASSO mixing parameter.  $\text{gamma\_w} = 1$  is LASSO and  $\text{gamma\_w} = 0$  group LASSO.  
 full\_est pre-estimated parameters based on full sample and `regress_choice` for a sequence of  $\lambda$ 's.

method_choice	choose between ‘initial’, ‘ic’ and ‘cv’. ‘initial’ pre-computes initial estimates. ‘ic’ computes solution based on information criteria (BIC, AIC or AICc). ‘cv’ computes solution based on cross-validation (cv).
nlam	number of $\lambda$ ’s to use in the regularization path.
lambdas	user specified sequence of $\lambda$ values for fitting. We recommend leaving this to NULL and letting function to self-select values.
min_frac	the minimum value of the penalty parameter, as a fraction of the maximum value.
nfolds	number of folds of the cv loop.
lambda_choice	choose between ‘min’ and ‘1se’. ‘min’ computes solution that minimizes the cv error. ‘1se’ computes solution such that the cv error is within 1 standard error of the minimum ‘min’.
ic_choice	choose between ‘bic’, ‘aic’ and ‘aicc’. ‘bic’ computes solution that minimizes Bayesian information criterion. ‘aic’ computes solution that minimizes Akaike information criterion. ‘aicc’ computes solution that minimizes Akaike information corrected criterion.
num_cores	number of cores used to compute cv loop.
verbose	flag to print information.
thresh	convergence threshold for change in beta. We recommend leaving this to NULL.
outer_thresh	outer loop convergence threshold. We recommend leaving this to NULL.
inner_iter	the maximum number of inner sg-LASSO loop iterations. We recommend leaving this to NULL.
outer_iter	the maximum number of outer sg-LASSO loop iterations. We recommend leaving this to NULL.

## Details

The sequence of linear regression models implied by lambdas vector is fit by block coordinate-descent. The objective function is

$$RSS(\alpha, \beta)/T + 2\lambda * \Omega_\gamma(\beta),$$

where  $RSS(\alpha, \beta)$  is the least squares fit. The penalty function  $\Omega_\gamma(\cdot)$  is applied on  $\beta$  coefficients and is

$$\Omega_\gamma(\beta) = \gamma|\beta|_1 + (1 - \gamma)\|\beta\|_{2,1},$$

a convex combination of LASSO and group LASSO penalty functions.

## Value

Parameter estimates of linear regression model under sg-LASSO penalty.

## Author(s)

Jonas Striaukas

## Examples

```
set.seed(1)
x = matrix(rnorm(100 * 20), 100, 20)
y = rnorm(100)
index = 1:20
reg_sgl(X = x, y = y, index = index, gamma_w = 1, method_choice = "initial",
        num_cores = 2, verbose = FALSE, lambdas = c(1,2,3))
```

*sgl\_fit*

*sg-LASSO regression*

## Description

Fits mse sg-LASSO regression model.

## Usage

```
sgl_fit(X, Z, y, index, lambdas, gamma_w = NULL, l1_factor = NULL, l21_factor = NULL,
        dummies_index = NULL, inner_iter = NULL, outer_iter = NULL, thresh = NULL,
        outer_thresh = NULL)
```

## Arguments

X	T by p data matrix, where t and p respectively denote the sample size and the number of regressors.
Z	dummies matrix.
y	T by 1 vector of outcome.
index	p by 1 vector indicating group membership of each covariate.
lambdas	user specified sequence of $\lambda$ values for fitting. We recommend leaving this to NULL and letting function to self-select values.
gamma_w	sg-LASSO mixing parameter. $\text{gamma\_w} = 1$ is LASSO and $\text{gamma\_w} = 0$ group LASSO.
l1_factor	$\ell_1$ norm penalty factor for random or fixed effects (default value is zero which means $\alpha$ is left unpenalized in $\ell_1$ norm).
l21_factor	$\ell_1$ norm penalty factor for random or fixed effects (default value is zero which means $\alpha$ is left unpenalized in $\ell_1$ norm).
dummies_index	vector indicating group membership of $\alpha$ (default - no grouping).
inner_iter	the maximum number of inner sg-LASSO loop iterations. We recommend leaving this to NULL.
outer_iter	the maximum number of outer sg-LASSO loop iterations. We recommend leaving this to NULL.
thresh	convergence threshold for change in beta. We recommend leaving this to NULL.
outer_thresh	outer loop convergence threshold. We recommend leaving this to NULL.

## Details

The sequence of linear regression models implied by `lambdas` vector is fit by block coordinate-descent. The objective function is

$$RSS(\alpha, \beta)/T + 2\lambda * \Omega_\gamma(\beta),$$

where  $RSS(\alpha, \beta)$  is the least squares fit. The penalty function  $\Omega_\gamma(\cdot)$  is applied on  $\beta$  coefficients and is

$$\Omega_\gamma(\beta) = \gamma|\beta|_1 + (1 - \gamma)\|\beta\|_{2,1},$$

a convex combination of LASSO and group LASSO penalty functions.

## Value

sg-LASSO regression fitted coefficients.

## Author(s)

Jonas Striaukas

## Examples

```
set.seed(1)
x = matrix(rnorm(100 * 20), 100, 20)
y = rnorm(100)
index = 1:20
Z <- as.matrix(rep(1,times=length(y)))
sgl_fit(X = x, Z = Z, y = y, index = index, lambdas = c(1,2,3), gamma_w = 1)
```

`transform_dt`

*Time series vector transformation*

## Description

Transform time series a vector based on transformation code.

## Usage

```
transform_dt(x, tcode)
```

## Arguments

<code>x</code>	time series vector.
<code>tcode</code>	transformation code. 1 - not transformed, 2 - first difference, 3 - second difference, 4 - natural log, 5 - first difference of natural log, 6 - second difference of natural log, 7 - first difference of percent change.

**Value**

transformed time series vector.

**Examples**

```
x <- rnorm(100,1)
transform_dt(x, tcode = 1)
```

*us\_rgdp*

*US real GDP data with several high-frequency predictors*

**Description**

US real GDP, Chicago National Activity Index, Nonfarm payrolls and ADS Index

**Usage**

```
data(us_rgdp)
```

**Format**

A `data.frame` object.a

**Source**

rgdp - [FRED](#)  
 cfnai - [Chicago Fed website](#)  
 payems - [FRED](#)  
 ads - [Philadelphia Fed website](#)

**Examples**

```
data(us_rgdp)
us_rgdp$rgdp # - GDP data
us_rgdp$cfnai # - CNAI predictor data
us_rgdp$payems # - Nonfarm payrolls predictor data
us_rgdp$ads # - ADS predictor data
```

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