

Package ‘dfms’

August 20, 2025

Version 0.3.1

Title Dynamic Factor Models

Description Efficient estimation of Dynamic Factor Models using the Expectation Maximization (EM) algorithm or Two-Step (2S) estimation, supporting datasets with missing data. Factors are assumed to follow a stationary VAR process of order p . The estimation options follow advances in the econometric literature: either running the Kalman Filter and Smoother once with initial values from PCA - 2S estimation as in Doz, Giannone and Reichlin (2011) <doi:10.1016/j.jeconom.2011.02.012> - or via iterated Kalman Filtering and Smoothing until EM convergence - following Doz, Giannone and Reichlin (2012) <doi:10.1162/REST_a_00225> - or using the adapted EM algorithm of Banbura and Modugno (2014) <doi:10.1002/jae.2306>, allowing arbitrary patterns of missing data. The implementation makes heavy use of the 'Armadillo' 'C++' library and the 'collapse' package, providing for particularly speedy estimation. A comprehensive set of methods supports interpretation and visualization of the model as well as forecasting. Information criteria to choose the number of factors are also provided - following Bai and Ng (2002) <doi:10.1111/1468-0262.00273>.

URL <https://sebkranztz.github.io/dfms/>,
<https://github.com/SebKranztz/dfms>

BugReports <https://github.com/SebKranztz/dfms/issues>

Depends R (>= 3.5.0)

Imports Rcpp (>= 1.0.1), collapse (>= 2.0.0)

LinkingTo Rcpp, RcppArmadillo

Suggests xts, vars, magrittr, testthat (>= 3.0.0), knitr, rmarkdown,
covr

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Encoding UTF-8
LazyData true
RoxygenNote 7.2.3
Config/testthat/edition 3
VignetteBuilder knitr
NeedsCompilation yes
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Repository CRAN
Date/Publication 2025-08-20 16:40:02 UTC

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Description

dfms provides efficient estimation of Dynamic Factor Models via the EM Algorithm — following Doz, Giannone & Reichlin (2011, 2012) and Banbura & Modugno (2014). Contents:

Information Criteria to Determine the Number of Factors

`ICr()`

- `plot(<ICr>)`
- `screeplot(<ICr>)`

Fit a Dynamic Factor Model

`DFM()`

- `summary(<dfm>)`
- `plot(<dfm>)`
- `as.data.frame(<dfm>)`
- `residuals(<dfm>)`
- `fitted(<dfm>)`

Generate Forecasts

`predict(<dfm>)`

- `plot(<dfm_forecast>)`
- `as.data.frame(<dfm_forecast>)`

Fast Stationary Kalman Filtering and Smoothing

`SKF()` — Stationary Kalman Filter

`FIS()` — Fixed Interval Smoother

`SKFS()` — Stationary Kalman Filter + Smoother

Helper Functions

`.VAR()` — (Fast) Barebones Vector-Autoregression

`ainv()` — Armadillo's Inverse Function

`apinv()` — Armadillo's Pseudo-Inverse Function

`tsnarmimp()` — Remove and Impute Missing Values in a Multivariate Time Series

`em_converged()` — Convergence Test for EM-Algorithm

Data

`BM14_M` — Monthly Series by Banbura and Modugno (2014)

`BM14_Q` — Quarterly Series by Banbura and Modugno (2014)

`BM14_Models` — Series Metadata + Small/Medium/Large Model Specifications

References

Doz, C., Giannone, D., & Reichlin, L. (2011). A two-step estimator for large approximate dynamic factor models based on Kalman filtering. *Journal of Econometrics*, 164(1), 188-205. doi:10.1016/j.jeconom.2011.02.012

Doz, C., Giannone, D., & Reichlin, L. (2012). A quasi-maximum likelihood approach for large, approximate dynamic factor models. *Review of Economics and Statistics*, 94(4), 1014-1024. doi:10.1162/REST_a_00225

Banbura, M., & Modugno, M. (2014). Maximum likelihood estimation of factor models on datasets with arbitrary pattern of missing data. *Journal of Applied Econometrics*, 29(1), 133-160. doi:10.1002/jae.2306

.VAR

(Fast) Barebones Vector-Autoregression

Description

Quickly estimate a VAR(p) model using Armadillo's inverse function.

Usage

```
.VAR(x, p = 1L)
```

Arguments

`x` data numeric matrix with time series in columns - without missing values.
`p` positive integer. The lag order of the VAR.

Value

A list containing matrices $Y = x[-(1:p),]$, X which contains lags 1 - p of x combined column-wise, A which is the $np \times n$ transition matrix, where n is the number of series in x , and the VAR residual matrix $res = Y - X \%*\% A$.

A list with the following elements:

`Y` $x[-(1:p),]$.
`X` lags 1 - p of x combined column-wise.
`A` $np \times n$ transition matrix, where n is the number of series in x .
`res` VAR residual matrix: $Y - X \%*\% A$.

See Also[dfms-package](#)**Examples**

```
var = .VAR(diff(EuStockMarkets), 3)
str(var)
var$A
rm(var)
```

ainv

Armadillo's Inverse Functions

Description

Matrix inverse and pseudo-inverse by the Armadillo C++ library.

Usage

```
ainv(x)
```

```
apinv(x)
```

Arguments

x a numeric matrix, must be square for ainv.

Value

The matrix-inverse or pseudo-inverse.

See Also[dfms-package](#)**Examples**

```
ainv(crossprod(diff(EuStockMarkets)))
```

as.data.frame.dfm *Extract Factor Estimates in a Data Frame*

Description

Extract Factor Estimates in a Data Frame

Usage

```
## S3 method for class 'dfm'
as.data.frame(
  x,
  ...,
  method = "all",
  pivot = c("long", "wide.factor", "wide.method", "wide", "t.wide"),
  time = seq_row(x$F_pca),
  stringsAsFactors = TRUE
)
```

Arguments

x	an object class 'dfm'.
...	not used.
method	character. The factor estimates to use: any of "qml", "2s", "pca" (multiple can be supplied) or "all" for all estimates.
pivot	character. The orientation of the frame: "long", "wide.factor" or "wide.method", "wide" or "t.wide".
time	a vector identifying the time dimension, or NULL to omit a time variable.
stringsAsFactors	make factors from method and factor identifiers. Same as option to as.data.frame.table .

Value

A data frame of factor estimates.

See Also

[dfms-package](#)

Examples

```
library(xts)
# Fit DFM with 3 factors and 3 lags in the transition equation
mod <- DFM(diff(BM14_M), r = 3, p = 3)

# Taking a single estimate:
print(head(as.data.frame(mod, method = "qml")))
```

```

print(head(as.data.frame(mod, method = "qml", pivot = "wide")))

# Adding a proper time variable
time <- index(BM14_M)[-1L]
print(head(as.data.frame(mod, method = "qml", time = time)))

# All estimates: different pivoting methods
for (pv in c("long", "wide.factor", "wide.method", "wide", "t.wide")) {
  cat("\npivot = ", pv, "\n")
  print(head(as.data.frame(mod, pivot = pv, time = time), 3))
}

```

BM14_Models

Euro Area Macroeconomic Data from Banbura and Modugno 2014

Description

A data extract from BM 2014 replication files. Some proprietary series (mostly PMI's) are excluded. The dataset BM14_Models provides information about all series and their inclusion in the 'small', 'medium' and 'large' sized dynamic factor models estimated by BM 2014. The actual data is contained in *xts* format in BM14_M for monthly data and BM14_Q for quarterly data.

Usage

```

BM14_Models
BM14_M
BM14_Q

```

Format

BM14_Models is a data frame with 101 obs. (series) and 8 columns:

series BM14 series code (converted to snake case for R)

label BM14 series label

code original series code from data source

freq series frequency

log_trans logical indicating whether the series was transformed by the natural log before differencing. Note that all data are provided in untransformed levels, and all data was (log-)differenced by BM14 before estimation.

small logical indicating series included in the 'small' model of BM14. Proprietary series are excluded.

medium logical indicating series included in the 'medium' model of BM14. Proprietary series are excluded.

large logical indicating series included in the 'large' model of BM14. This comprises all series, thus the variable is redundant but included for completeness. Proprietary series are excluded.

Source

Banbura, M., & Modugno, M. (2014). Maximum likelihood estimation of factor models on datasets with arbitrary pattern of missing data. *Journal of Applied Econometrics*, 29(1), 133-160.

See Also

[dfms-package](#)

Examples

```
library(magrittr)
library(xts)

# Constructing the database for the large model
BM14 = merge(BM14_M, BM14_Q)
BM14[, BM14_Models$log_trans] %<>% log()
BM14[, BM14_Models$freq == "M"] %<>% diff()
BM14[, BM14_Models$freq == "Q"] %<>% diff(3)

# Small Model Database
head(BM14[, BM14_Models$small])

# Medium-Sized Model Database
head(BM14[, BM14_Models$medium])
```

DFM

Estimate a Dynamic Factor Model

Description

Efficient estimation of a Dynamic Factor Model via the EM Algorithm - on stationary data with time-invariant system matrices and classical assumptions, while permitting missing data.

Usage

```
DFM(
  X,
  r,
  p = 1L,
  ...,
  idio.ar1 = FALSE,
  quarterly.vars = NULL,
  rQ = c("none", "diagonal", "identity"),
  rR = c("diagonal", "identity", "none"),
  em.method = c("auto", "DGR", "BM", "none"),
  min.iter = 25L,
  max.iter = 100L,
```

```

    tol = 1e-04,
    pos.corr = TRUE,
    check.increased = FALSE
  )

```

Arguments

<code>X</code>	a $T \times n$ numeric data matrix or frame of stationary time series. May contain missing values. <i>Note</i> that data is internally standardized (scaled and centered) before estimation.
<code>r</code>	integer. Number of factors.
<code>p</code>	integer. Number of lags in factor VAR.
<code>...</code>	(optional) arguments to tsnarmimp . The default settings impute internal missing values with a cubic spline and the edges with the median and a 3-period moving average.
<code>idio.ar1</code>	logical. Model observation errors as AR(1) processes: $e_t = \rho e_{t-1} + v_t$. <i>Note</i> that this substantially increases computation time, and is generally not needed if n is large (>30). See theoretical vignette for details.
<code>quarterly.vars</code>	character. Names of quarterly variables in <code>X</code> (if any). Monthly variables should be to the left of the quarterly variables in the data matrix and quarterly observations should be provided every 3rd period.
<code>rQ</code>	character. Restrictions on the state (transition) covariance matrix (Q).
<code>rR</code>	character. Restrictions on the observation (measurement) covariance matrix (R).
<code>em.method</code>	character. The implementation of the Expectation Maximization Algorithm used. The options are:
<code>"auto"</code>	Automatic selection: "BM" if anyNA(<code>X</code>), else "DGR".
<code>"DGR"</code>	The classical EM implementation of Doz, Giannone and Reichlin (2012). This implementation is efficient and quick.
<code>"BM"</code>	The modified EM algorithm of Banbura and Modugno (2014) which also accounts for missing data in the EM iterations.
<code>"none"</code>	Performs no EM iterations and just returns the Two-Step estimates from running the data through the Kalman Filter.
<code>min.iter</code>	integer. Minimum number of EM iterations (to ensure a convergence path).
<code>max.iter</code>	integer. Maximum number of EM iterations.
<code>tol</code>	numeric. EM convergence tolerance.
<code>pos.corr</code>	logical. Increase the likelihood that factors correlate positively with the data, by scaling the eigenvectors such that the principal components (used to initialize the Kalman Filter) co-vary positively with the row-means of the standardized data.
<code>check.increased</code>	logical. Check if likelihood has increased. Passed to em_converged . If TRUE, the algorithm only terminates if convergence was reached with decreasing likelihood.

Details

This function efficiently estimates a Dynamic Factor Model with the following classical assumptions:

1. Linearity
2. Idiosyncratic measurement (observation) errors (R is diagonal)
3. No direct relationship between series and lagged factors (*ceteris paribus* contemporaneous factors)
4. No relationship between lagged error terms in the either measurement or transition equation (no serial correlation), unless explicitly modeled as AR(1) processes using `idio.ar1 = TRUE`.

Factors are allowed to evolve in a $VAR(p)$ process, and data is internally standardized (scaled and centered) before estimation (removing the need of intercept terms). By assumptions 1-4, this translates into the following dynamic form:

$$\begin{aligned}\mathbf{x}_t &= \mathbf{C}_0 \mathbf{f}_t + \mathbf{e}_t \sim N(\mathbf{0}, \mathbf{R}) \\ \mathbf{f}_t &= \sum_{j=1}^p \mathbf{A}_j \mathbf{f}_{t-j} + \mathbf{u}_t \sim N(\mathbf{0}, \mathbf{Q}_0)\end{aligned}$$

where the first equation is called the measurement or observation equation and the second equation is called transition, state or process equation, and

n	number of series in $\mathbf{x}_t$ (r and p as the arguments to DFM).
$\mathbf{x}_t$	$n \times 1$ vector of observed series at time t : $(x_{1t}, \dots, x_{nt})'$. Some observations can be missing.
$\mathbf{f}_t$	$r \times 1$ vector of factors at time t : $(f_{1t}, \dots, f_{rt})'$.
$\mathbf{C}_0$	$n \times r$ measurement (observation) matrix.
$\mathbf{A}_j$	$r \times r$ state transition matrix at lag j .
$\mathbf{Q}_0$	$r \times r$ state covariance matrix.
$\mathbf{R}$	$n \times n$ measurement (observation) covariance matrix. It is diagonal by assumption 2 that $E[\mathbf{x}_{it} \mathbf{x}_{-i,t}, \mathbf{x}_{i,t-1}, \dots, \mathbf{f}_t, \mathbf{f}_t]$

This model can be estimated using a classical form of the Kalman Filter and the Expectation Maximization (EM) algorithm, after transforming it to State-Space (stacked, VAR(1)) form:

$$\begin{aligned}\mathbf{x}_t &= \mathbf{C} \mathbf{F}_t + \mathbf{e}_t \sim N(\mathbf{0}, \mathbf{R}) \\ \mathbf{F}_t &= \mathbf{A} \mathbf{F}_{t-1} + \mathbf{u}_t \sim N(\mathbf{0}, \mathbf{Q})\end{aligned}$$

where

n	number of series in $\mathbf{x}_t$ (r and p as the arguments to DFM).
$\mathbf{x}_t$	$n \times 1$ vector of observed series at time t : $(x_{1t}, \dots, x_{nt})'$. Some observations can be missing.
$\mathbf{F}_t$	$rp \times 1$ vector of stacked factors at time t : $(f_{1t}, \dots, f_{rt}, f_{1,t-1}, \dots, f_{r,t-1}, \dots, f_{1,t-p}, \dots, f_{r,t-p})'$.
$\mathbf{C}$	$n \times rp$ observation matrix. Only the first $n \times r$ terms are non-zero, by assumption 3 that $E[\mathbf{x}_t \mathbf{F}_t] = E[\mathbf{x}_t \mathbf{f}_t]$ (no relationship between factors at different times).
$\mathbf{A}$	stacked $rp \times rp$ state transition matrix consisting of 3 parts: the top $r \times rp$ part provides the dynamic relationships across time, the bottom left $p \times r$ part provides the dynamic relationships across factors, and the bottom right $p \times p$ part provides the dynamic relationships across lags.
$\mathbf{Q}$	$rp \times rp$ state covariance matrix. The top $r \times r$ part gives the contemporaneous relationships, the rest are zeros by assumption 4.
$\mathbf{R}$	$n \times n$ observation covariance matrix. It is diagonal by assumption 2 and identical to $\mathbf{C}$ as stated in the dynamic form.

The filter is initialized with PCA estimates on the imputed dataset—see [SKFS](#) for a complete code example.

Value

A list-like object of class 'dfm' with the following elements:

<code>X_imp</code>	$T \times n$ matrix with the imputed and standardized (scaled and centered) data—after applying tsnarmimp . It has attributes attached allowing for reconstruction of the original data:
<code>"stats"</code>	is a $n \times 5$ matrix of summary statistics of class "qsu" (see qsu).
<code>"missing"</code>	is a $T \times n$ logical matrix indicating missing or infinite values in the original data (which are imputed in <code>X_imp</code>).
<code>"attributes"</code>	contains the attributes of the original data input.
<code>"is.list"</code>	is a logical value indicating whether the original data input was a list / data frame.
<code>eigen</code>	<code>eigen(cov(X_imp))</code> .
<code>F_pca</code>	$T \times r$ matrix of principal component factor estimates - <code>X_imp %%% eigen\$vector</code> s.
<code>P_0</code>	$r \times r$ initial factor covariance matrix estimate based on PCA results.
<code>F_2s</code>	$T \times r$ matrix two-step factor estimates as in Doz, Giannone and Reichlin (2011) - obtained from running the data through the Kalman Filter and Smoother once, where the Filter is initialized with results from PCA.
<code>P_2s</code>	$r \times r \times T$ covariance matrices of two-step factor estimates.
<code>F_qml</code>	$T \times r$ matrix of quasi-maximum likelihood factor estimates - obtained by iteratively Kalman Filtering and Smoothing the factor estimates until EM convergence.

P_qml	$r \times r \times T$ covariance matrices of QML factor estimates.
A	$r \times rp$ factor transition matrix.
C	$n \times r$ observation matrix.
Q	$r \times r$ state (error) covariance matrix.
R	$n \times n$ observation (error) covariance matrix.
e	$T \times n$ estimates of observation errors $\mathbf{e}_t$. Only available if <code>idio.ar1 = TRUE</code> .
rho	$n \times 1$ estimates of AR(1) coefficients (ρ) in observation errors: $e_t = \rho e_{t-1} + v_t$. Only available if <code>idio.ar1 = TRUE</code> .
loglik	vector of log-likelihoods - one for each EM iteration. The final value corresponds to the log-likelihood of the reported model.
tol	The numeric convergence tolerance used.
converged	single logical valued indicating whether the EM algorithm converged (within <code>max.iter</code> iterations subject to <code>tol</code>).
anyNA	single logical valued indicating whether there were any (internal) missing values in the data (determined after removal of rows with too many missing values). If FALSE, <code>X_imp</code> is simply the original data in matrix form, and does not have the "missing" attribute attached.
rm.rows	vector of any cases (rows) that were removed beforehand (subject to <code>max.missing</code> and <code>na.rm.method</code>). If no cases were removed the slot is NULL.
quarterly.vars	names of the quarterly variables (if any).
em.method	The EM method used.
call	call object obtained from <code>match.call()</code> .

References

- Doz, C., Giannone, D., & Reichlin, L. (2011). A two-step estimator for large approximate dynamic factor models based on Kalman filtering. *Journal of Econometrics*, 164(1), 188-205.
- Doz, C., Giannone, D., & Reichlin, L. (2012). A quasi-maximum likelihood approach for large, approximate dynamic factor models. *Review of Economics and Statistics*, 94(4), 1014-1024.
- Banbura, M., & Modugno, M. (2014). Maximum likelihood estimation of factor models on datasets with arbitrary pattern of missing data. *Journal of Applied Econometrics*, 29(1), 133-160.
- Stock, J. H., & Watson, M. W. (2016). Dynamic Factor Models, Factor-Augmented Vector Autoregressions, and Structural Vector Autoregressions in Macroeconomics. *Handbook of Macroeconomics*, 2, 415–525. <https://doi.org/10.1016/bs.hesmac.2016.04.002>

See Also

[dfms-package](#)

Examples

```

library(magrittr)
library(xts)
library(vars)

# BM14 Replication Data. Constructing the database:
BM14 <- merge(BM14_M, BM14_Q)
BM14[, BM14_Models$log_trans] %<>% log()
BM14[, BM14_Models$freq == "M"] %<>% diff()
BM14[, BM14_Models$freq == "Q"] %<>% diff(3)

### Small Model -----

# IC for number of factors
IC_small <- ICr(BM14[, BM14_Models$small], max.r = 5)
plot(IC_small)
screplot(IC_small)

# I take 2 factors. Now number of lags
VARselect(IC_small$F_pca[, 1:2])

# Estimating the model with 2 factors and 3 lags
dfm_small <- DFM(BM14[, BM14_Models$small], r = 2, p = 3,
  quarterly.vars = BM14_Models %$% series[freq == "Q" & small])

# Inspecting the model
summary(dfm_small)
plot(dfm_small) # Factors and data
plot(dfm_small, method = "all", type = "individual") # Factor estimates
plot(dfm_small, type = "residual") # Residuals from factor predictions

# 10 periods ahead forecast
plot(predict(dfm_small), xlim = c(300, 370))

### Medium-Sized Model -----

# IC for number of factors
IC_medium <- ICr(BM14[, BM14_Models$medium])
plot(IC_medium)
screplot(IC_medium)

# I take 3 factors. Now number of lags
VARselect(IC_medium$F_pca[, 1:3])

# Estimating the model with 3 factors and 3 lags
dfm_medium <- DFM(BM14[, BM14_Models$medium], r = 3, p = 3,
  quarterly.vars = BM14_Models %$% series[freq == "Q" & medium])

# Inspecting the model
summary(dfm_medium)

```

```

plot(dfm_medium) # Factors and data
plot(dfm_medium, method = "all", type = "individual") # Factor estimates
plot(dfm_medium, type = "residual") # Residuals from factor predictions

# 10 periods ahead forecast
plot(predict(dfm_medium), xlim = c(300, 370))

### Large Model -----

# IC for number of factors
IC_large <- ICr(BM14)
plot(IC_large)
screeplot(IC_large)

# I take 6 factors. Now number of lags
VARselect(IC_large$F_pca[, 1:6])

# Estimating the model with 6 factors and 3 lags
dfm_large <- DFM(BM14, r = 6, p = 3,
  quarterly.vars = BM14_Models %%% series[freq == "Q"])

# Inspecting the model
summary(dfm_large)
plot(dfm_large) # Factors and data
# plot(dfm_large, method = "all", type = "individual") # Factor estimates
plot(dfm_large, type = "residual") # Residuals from factor predictions

# 10 periods ahead forecast
plot(predict(dfm_large), xlim = c(300, 370))

```

em_converged

Convergence Test for EM-Algorithm

Description

Convergence Test for EM-Algorithm

Usage

```
em_converged(loglik, previous_loglik, tol = 1e-04, check.increased = FALSE)
```

Arguments

loglik	numeric. Current value of the log-likelihood function.
previous_loglik	numeric. Value of the log-likelihood function at the previous iteration.
tol	numerical. The tolerance of the test. If $ LL(t) - LL(t-1) / \text{avg} < \text{tol}$, where $\text{avg} = (LL(t) + LL(t-1))/2$, then algorithm has converged.

`check.increased`
 logical. Check if likelihood has increased.

Value

A logical statement indicating whether EM algorithm has converged. if `check.increased = TRUE`, a vector with 2 elements indicating the convergence status and whether the likelihood has decreased.

See Also

[dfms-package](#)

Examples

```
em_converged(1001, 1000)
em_converged(10001, 10000)
em_converged(10001, 10000, check = TRUE)
em_converged(10000, 10001, check = TRUE)
```

FIS

(Fast) Fixed-Interval Smoother (Kalman Smoother)

Description

(Fast) Fixed-Interval Smoother (Kalman Smoother)

Usage

```
FIS(A, F, F_pred, P, P_pred, F_0 = NULL, P_0 = NULL)
```

Arguments

A	transition matrix ($rp \times rp$).
F	state estimates ($T \times rp$).
F_pred	state predicted estimates ($T \times rp$).
P	variance estimates ($rp \times rp \times T$).
P_pred	predicted variance estimates ($rp \times rp \times T$).
F_0	initial state vector ($rp \times 1$) or empty (NULL).
P_0	initial state covariance ($rp \times rp$) or empty (NULL).

Details

The Kalman Smoother is given by:

$$\begin{aligned}\mathbf{J}_t &= \mathbf{P}_t \mathbf{A} + inv(\mathbf{P}_{t+1}^{pred}) \\ \mathbf{F}_t^{smooth} &= \mathbf{F}_t + \mathbf{J}_t(\mathbf{F}_{t+1}^{smooth} - \mathbf{F}_{t+1}^{pred}) \\ \mathbf{P}_t^{smooth} &= \mathbf{P}_t + \mathbf{J}_t(\mathbf{P}_{t+1}^{smooth} - \mathbf{P}_{t+1}^{pred})\mathbf{J}_t'\end{aligned}$$

The initial smoothed values for period $t = T$ are set equal to the filtered values. If $\mathbf{F}_0$ and $\mathbf{P}_0$ are supplied, the smoothed initial conditions ($t = 0$ values) are also calculated and returned. For further details see any textbook on time series such as Shumway & Stoffer (2017), which provide an analogous R implementation in `astsa::Ksmooth0`.

Value

Smoothed state and covariance estimates, including initial ($t = 0$) values.

<code>F_smooth</code>	$T \times rp$ smoothed state vectors, equal to the filtered state in period T .
<code>P_smooth</code>	$rp \times rp \times T$ smoothed state covariance, equal to the filtered covariance in period T .
<code>F_smooth_0</code>	$1 \times rp$ initial smoothed state vectors, based on $\mathbf{F}_0$.
<code>P_smooth_0</code>	$rp \times rp$ initial smoothed state covariance, based on $\mathbf{P}_0$.

References

- Shumway, R. H., & Stoffer, D. S. (2017). Time Series Analysis and Its Applications: With R Examples. Springer.
- Harvey, A. C. (1990). Forecasting, structural time series models and the Kalman filter.

See Also

[SKF SKFS dfms-package](#)

Examples

```
# See ?SKFS
```

ICr	<i>Information Criteria to Determine the Number of Factors (r)</i>
-----	--

Description

Minimizes 3 information criteria proposed by Bai and Ng (2002) to determine the optimal number of factors r^* to be used in an approximate factor model. A Screeplot can also be computed to eyeball the number of factors in the spirit of Onatski (2010).

Usage

```
ICr(X, max.r = min(20, ncol(X) - 1))

## S3 method for class 'ICr'
print(x, ...)

## S3 method for class 'ICr'
plot(x, ...)

## S3 method for class 'ICr'
screeplot(x, type = "pve", show.grid = TRUE, max.r = 30, ...)
```

Arguments

X	a $T \times n$ numeric data matrix or frame of stationary time series.
max.r	integer. The maximum number of factors for which IC should be computed (or eigenvalues to be displayed in the screeplot).
x	an object of type 'ICr'.
...	further arguments to ts.plot or plot .
type	character. Either "ev" (eigenvalues), "pve" (percent variance explained), or "cum.pve" (cumulative PVE). Multiple plots can be requested.
show.grid	logical. TRUE shows gridlines in each plot.

Details

Following Bai and Ng (2002) and De Valk et al. (2019), let $NSSR(r)$ be the normalized sum of squared residuals $SSR(r)/(n \times T)$ when r factors are estimated using principal components. Then the information criteria can be written as follows:

$$IC_{r1} = \ln(NSSR(r)) + r \left(\frac{n+T}{nT} \right) + \ln \left(\frac{nT}{n+T} \right)$$

$$IC_{r2} = \ln(NSSR(r)) + r \left(\frac{n+T}{nT} \right) + \ln(\min(n, T))$$

$$IC_{r3} = \ln(NSSR(r)) + r \left(\frac{\ln(\min(n, T))}{\min(n, T)} \right)$$

The optimal number of factors r^* corresponds to the minimum IC. The three criteria are asymptotically equivalent, but may give significantly different results for finite samples. The penalty in $IC_{r,2}$ is highest in finite samples.

In the Screeplot a horizontal dashed line is shown signifying an eigenvalue of 1, or a share of variance corresponding to 1 divided by the number of eigenvalues.

Value

A list of 4 elements:

<code>F_pca</code>	T x n matrix of principle component factor estimates.
<code>eigenvalues</code>	the eigenvalues of the covariance matrix of X.
<code>IC</code>	<code>r.max</code> x 3 'table' containing the 3 information criteria of Bai and Ng (2002), computed for all values of <code>r</code> from <code>1:r.max</code> .
<code>r.star</code>	vector of length 3 containing the number of factors (<code>r</code>) minimizing each information criterion.

Note

To determine the number of lags (`p`) in the factor transition equation, use the function `vars::VARselect` with `r*` principle components (also returned by `ICr`).

References

- Bai, J., Ng, S. (2002). Determining the Number of Factors in Approximate Factor Models. *Econometrica*, 70(1), 191-221. <https://doi.org/10.1111/1468-0262.00273>.
- Onatski, A. (2010). Determining the number of factors from empirical distribution of eigenvalues. *The Review of Economics and Statistics*, 92(4), 1004-1016.
- De Valk, S., de Mattos, D., & Ferreira, P. (2019). Nowcasting: An R package for predicting economic variables using dynamic factor models. *The R Journal*, 11(1), 230-244.

See Also

[dfms-package](#)

Examples

```
library(xts)
library(vars)

ics <- ICr(diff(BM14_M))
print(ics)
plot(ics)
screeplot(ics)

# Optimal lag-order with 6 factors chosen
VARselect(ics$F_pca[, 1:6])
```

plot.dfm

*Plot DFM***Description**

Plot DFM

Usage

```
## S3 method for class 'dfm'
plot(
  x,
  method = switch(x$em.method, none = "2s", "qml"),
  type = c("joint", "individual", "residual"),
  scale.factors = TRUE,
  ...
)

## S3 method for class 'dfm'
screepplot(x, ...)
```

Arguments

x	an object class 'dfm'.
method	character. The factor estimates to use: one of "qml", "2s", "pca" or "all" to plot all estimates.
type	character. The type of plot: "joint", "individual" or "residual".
scale.factors	logical. Standardize factor estimates, this usually improves the plot since the factor estimates corresponding to the greatest PCA eigenvalues tend to have a greater variance than the data.
...	for plot.dfm: further arguments to plot , ts.plot , or boxplot , depending on the type of plot. For screepplot.dfm: further arguments to screepplot.ICr .

Value

Nothing.

See Also[dfms-package](#)**Examples**

```
# Fit DFM with 3 factors and 3 lags in the transition equation
mod <- DFM(diff(BM14_M), r = 3, p = 3)
plot(mod)
plot(mod, type = "individual", method = "all")
```

```
plot(mod, type = "residual")
```

predict.dfm

DFM Forecasts

Description

This function produces h-step ahead forecasts of both the factors and the data, with an option to also forecast autocorrelated residuals with a univariate method and produce a combined forecast.

Usage

```
## S3 method for class 'dfm'
predict(
  object,
  h = 10L,
  method = switch(object$em.method, none = "2s", "qml"),
  standardized = TRUE,
  resFUN = NULL,
  resAC = 0.1,
  ...
)
```

```
## S3 method for class 'dfm_forecast'
print(x, digits = 4L, ...)
```

```
## S3 method for class 'dfm_forecast'
plot(
  x,
  main = paste(x$h, "Period Ahead DFM Forecast"),
  xlab = "Time",
  ylab = "Standardized Data",
  factors = seq_len(ncol(x$F)),
  scale.factors = TRUE,
  factor.col = rainbow(length(factors)),
  factor.lwd = 1.5,
  fcst.lty = "dashed",
  data.col = c("grey85", "grey65"),
  legend = TRUE,
  legend.items = paste0("f", factors),
  grid = FALSE,
  vline = TRUE,
  vline.lty = "dotted",
  vline.col = "black",
  ...
)
```

```

)

## S3 method for class 'dfm_forecast'
as.data.frame(
  x,
  ...,
  use = c("factors", "data", "both"),
  pivot = c("long", "wide"),
  time = seq_len(nrow(x$F) + x$h),
  stringsAsFactors = TRUE
)

```

Arguments

object	an object of class 'dfm'.
h	integer. The forecast horizon.
method	character. The factor estimates to use: one of "qml", "2s" or "pca".
standardized	logical. FALSE will return data forecasts on the original scale.
resFUN	an (optional) function to compute a univariate forecast of the residuals. The function needs to have a second argument providing the forecast horizon (h) and return a vector of forecasts. See Examples.
resAC	numeric. Threshold for residual autocorrelation to apply resFUN: only residual series where $AC1 > resAC$ will be forecasted.
...	not used.
x	an object class 'dfm_forecast'.
digits	integer. The number of digits to print out.
main, xlab, ylab	character. Graphical parameters passed to ts.plot .
factors	integers indicating which factors to display. Setting this to NA, NULL or 0 will omit factor plots.
scale.factors	logical. Standardize factor estimates, this usually improves the plot since the factor estimates corresponding to the greatest PCA eigenvalues tend to have a greater variance than the data.
factor.col, factor.lwd	graphical parameters affecting the colour and line width of factor estimates plots. See par .
fcst.lty	integer or character giving the line type of the forecasts of factors and data. See par .
data.col	character vector of length 2 indicating the colours of historical data and forecasts of that data. Setting this to NA, NULL or "" will not plot data and data forecasts.
legend	logical. TRUE draws a legend in the top-left of the chart.
legend.items	character names of factors for the legend.
grid	logical. TRUE draws a grid on the background of the plot.
vline	logical. TRUE draws a vertical line delimitating historical data and forecasts.

<code>vline.lty, vline.col</code>	graphical parameters affecting the appearance of the vertical line. See par .
<code>use</code>	character. Which forecasts to use "factors", "data" or "both".
<code>pivot</code>	character. The orientation of the frame: "long" or "wide".
<code>time</code>	a vector identifying the time dimension, must be of length $T + h$, or NULL to omit a time variable.
<code>stringsAsFactors</code>	logical. If TRUE and <code>pivot = "long"</code> the 'Variable' column is created as a factor. Same as option to as.data.frame.table .

Value

A list-like object of class 'dfm_forecast' with the following elements:

<code>X_fcst</code>	$h \times n$ matrix with the forecasts of the variables.
<code>F_fcst</code>	$h \times r$ matrix with the factor forecasts.
<code>X</code>	$T \times n$ matrix with the standardized (scaled and centered) data - with attributes attached allowing reconstruction of the original data:
<code>"stats"</code>	is a $n \times 5$ matrix of summary statistics of class "qsu" (see qsu). Only attached if <code>standardized = TRUE</code> .
<code>"attributes"</code>	contains the attributes of the original data input.
<code>"is.list"</code>	is a logical value indicating whether the original data input was a list / data frame.
<code>F</code>	$T \times r$ matrix of factor estimates.
<code>method</code>	the factor estimation method used.
<code>anyNA</code>	logical indicating whether <code>X</code> contains any missing values.
<code>h</code>	the forecast horizon.
<code>resid.fc</code>	logical indicating whether a univariate forecasting function was applied to the residuals.
<code>resid.fc.ind</code>	indices indicating for which variables (columns of <code>X</code>) the residuals were forecasted using the univariate function.
<code>call</code>	call object obtained from <code>match.call()</code> .

See Also

[dfms-package](#)

Examples

```

library(xts)
library(collapse)

# Fit DFM with 3 factors and 3 lags in the transition equation
mod <- DFM(diff(BM14_M), r = 3, p = 3)

# 15 period ahead forecast
fc <- predict(mod, h = 15)
print(fc)
plot(fc, xlim = c(300, 370))

# Also forecasting autocorrelated residuals with an AR(1)
fcfun <- function(x, h) predict(ar(na_rm(x)), n.ahead = h)$pred
fcar <- predict(mod, resFUN = fcfun, h = 15)
plot(fcar, xlim = c(300, 370))

# Retrieving a data frame of the forecasts
head(as.data.frame(fcar, pivot = "wide")) # Factors
head(as.data.frame(fcar, use = "data"))    # Data
head(as.data.frame(fcar, use = "both"))    # Both

```

residuals.dfm

*DFM Residuals and Fitted Values***Description**

The residuals $\mathbf{e}_t = \mathbf{x}_t - \mathbf{CF}_t$ or fitted values $\mathbf{CF}_t$ of the DFM observation equation.

Usage

```

## S3 method for class 'dfm'
residuals(
  object,
  method = switch(object$em.method, none = "2s", "qml"),
  orig.format = FALSE,
  standardized = FALSE,
  na.keep = TRUE,
  ...
)

## S3 method for class 'dfm'
fitted(
  object,
  method = switch(object$em.method, none = "2s", "qml"),
  orig.format = FALSE,
  standardized = FALSE,

```

```

    na.keep = TRUE,
    ...
  )

```

Arguments

<code>object</code>	an object of class 'dfm'.
<code>method</code>	character. The factor estimates to use: one of "qml", "2s" or "pca".
<code>orig.format</code>	logical. TRUE returns residuals/fitted values in a data format similar to X.
<code>standardized</code>	logical. FALSE will put residuals/fitted values on the original data scale.
<code>na.keep</code>	logical. TRUE inserts missing values where X is missing (default TRUE as residuals/fitted values are only defined for observed data). FALSE returns the raw prediction, which can be used to interpolate data based on the DFM. For residuals, FALSE returns the difference between the prediction and the initial imputed version of X use for PCA to initialize the Kalman Filter.
<code>...</code>	not used.

Value

A matrix of DFM residuals or fitted values. If `orig.format = TRUE` the format may be different, e.g. a data frame.

See Also

[dfms-package](#)

Examples

```

library(xts)
# Fit DFM with 3 factors and 3 lags in the transition equation
mod <- DFM(diff(BM14_M), r = 3, p = 3)

# Residuals
head(resid(mod))
plot(resid(mod, orig.format = TRUE)) # this is an xts object

# Fitted values
head(fitted(mod))
head(fitted(mod, orig.format = TRUE)) # this is an xts object

```

SKF

*(Fast) Stationary Kalman Filter***Description**

A simple and fast C++ implementation of the Kalman Filter for stationary data (or random walks - data should be mean zero and without a trend) with time-invariant system matrices and missing data.

Usage

```
SKF(X, A, C, Q, R, F_0, P_0, loglik = FALSE)
```

Arguments

X	numeric data matrix ($T \times n$).
A	transition matrix ($rp \times rp$).
C	observation matrix ($n \times rp$).
Q	state covariance ($rp \times rp$).
R	observation covariance ($n \times n$).
F_0	initial state vector ($rp \times 1$).
P_0	initial state covariance ($rp \times rp$).
loglik	logical. Compute log-likelihood?

Details

The underlying state space model is:

$$\mathbf{x}_t = \mathbf{C}\mathbf{F}_t + \mathbf{e}_t \sim N(\mathbf{0}, \mathbf{R})$$

$$\mathbf{F}_t = \mathbf{A}\mathbf{F}_{t-1} + \mathbf{u}_t \sim N(\mathbf{0}, \mathbf{Q})$$

where x_t is $X[t,]$. The filter then first performs a time update (prediction)

$$\mathbf{F}_t = \mathbf{A}\mathbf{F}_{t-1}$$

$$\mathbf{P}_t = \mathbf{A}\mathbf{P}_{t-1}\mathbf{A}' + \mathbf{Q}$$

where $P_t = Cov(F_t)$. This is followed by the measurement update (filtering)

$$\mathbf{K}_t = \mathbf{P}_t\mathbf{C}'(\mathbf{C}\mathbf{P}_t\mathbf{C}' + \mathbf{R})^{-1}$$

$$\mathbf{F}_t = \mathbf{F}_t + \mathbf{K}_t(\mathbf{x}_t - \mathbf{C}\mathbf{F}_t)$$

$$\mathbf{P}_t = \mathbf{P}_t - \mathbf{K}_t\mathbf{C}\mathbf{P}_t$$

If a row of the data is all missing the measurement update is skipped i.e. the prediction becomes the filtered value. The log-likelihood is computed as

$$1/2 \sum_t \log(|St|) - e_t' S_t e_t - n \log(2\pi)$$

where $S_t = (CP_t C' + R)^{-1}$ and $e_t = x_t - CF_t$ is the prediction error.

For further details see any textbook on time series such as Shumway & Stoffer (2017), which provide an analogous R implementation in `astsa::Kfilter0`. For another fast (C-based) implementation that also allows time-varying system matrices and non-stationary data see `FKF::fkf`.

Value

Predicted and filtered state vectors and covariances.

F	$T \times rp$ filtered state vectors.
P	$rp \times rp \times T$ filtered state covariances.
F_pred	$T \times rp$ predicted state vectors.
P_pred	$rp \times rp \times T$ predicted state covariances.
loglik	value of the log likelihood.

References

- Shumway, R. H., & Stoffer, D. S. (2017). Time Series Analysis and Its Applications: With R Examples. Springer.
- Harvey, A. C. (1990). Forecasting, structural time series models and the Kalman filter.
- Hamilton, J. D. (1994). Time Series Analysis. Princeton university press.

See Also

[FIS SKFS dfms-package](#)

Examples

```
# See ?SKFS
```

SKFS

(Fast) Stationary Kalman Filter and Smoother

Description

(Fast) Stationary Kalman Filter and Smoother

Usage

```
SKFS(X, A, C, Q, R, F_0, P_0, loglik = FALSE)
```

Arguments

X	numeric data matrix ($T \times n$).
A	transition matrix ($rp \times rp$).
C	observation matrix ($n \times rp$).
Q	state covariance ($rp \times rp$).
R	observation covariance ($n \times n$).
F_0	initial state vector ($rp \times 1$).
P_0	initial state covariance ($rp \times rp$).
loglik	logical. Compute log-likelihood?

Value

All results from [SKF](#) and [FIS](#), and additionally a $rp \times rp \times T$ matrix PPM_smooth, which is equal to the estimate of $Cov(F_t^{smooth}, F_{t-1}^{smooth} | T)$ and needed for EM iterations. See 'Property 6.3: The Lag-One Covariance Smoother' in Shumway & Stoffer (2017).

References

Shumway, R. H., & Stoffer, D. S. (2017). Time Series Analysis and Its Applications: With R Examples. Springer.

See Also

[SKF FIS dfms-package](#)

Examples

```
library(collapse)

## Two-Step factor estimates from monthly BM (2014) data
X <- fscale(diff(qM(BM14_M))) # Standardizing as KF has no intercept
r <- 5L # 5 Factors
p <- 3L # 3 Lags
n <- ncol(X)

## Initializing the Kalman Filter with PCA results
X_imp <- tsnaimp(X) # Imputing Data
v <- eigen(cov(X_imp))$vectors[, 1:r] # PCA
F_pc <- X_imp %*% v # Principal component factor estimates
C <- cbind(v, matrix(0, n, r*p-r)) # Observation matrix
res <- X - tcrossprod(F_pc, v) # Residuals from static predictions
R <- diag(fvar(res)) # Observation residual covariance
var <- .VAR(F_pc, p) # VAR(p)
A <- rbind(t(var$A), diag(1, r*p-r, r*p))
Q <- matrix(0, r*p, r*p) # VAR residual matrix
Q[1:r, 1:r] <- cov(var$res)
F_0 <- var$X[1L, ] # Initial factor estimate and covariance
P_0 <- ainv(diag((r*p)^2) - kronecker(A,A)) %*% unattrib(Q)
```

```

dim(P_0) <- c(r*p, r*p)

## Run standartized data through Kalman Filter and Smoother once
kfs_res <- SKFS(X, A, C, Q, R, F_0, P_0, FALSE)

## Two-step solution is state mean from the Kalman Smoother
F_kal <- kfs_res$F_smooth[, 1:r, drop = FALSE]
colnames(F_kal) <- paste0("f", 1:r)

## See that this is equal to the Two-Step estimate by DFM()
all.equal(F_kal, DFM(X, r, p, em.method = "none", pos.corr = FALSE)$F_2s)

## Same in two steps using SKF() and FIS()
kfs_res2 <- with(SKF(X, A, C, Q, R, F_0, P_0, FALSE), FIS(A, F, F_pred, P, P_pred))
F_kal2 <- kfs_res2$F_smooth[, 1:r, drop = FALSE]
colnames(F_kal2) <- paste0("f", 1:r)
all.equal(F_kal, F_kal2)

rm(X, r, p, n, X_imp, v, F_pc, C, res, R, var, A, Q, F_0, P_0, kfs_res, F_kal, kfs_res2, F_kal2)

```

summary.dfm

DFM Summary Methods

Description

Summary and print methods for class 'dfm'. `print.dfm` just prints basic model information and the factor transition matrix **A**, `coef.dfm` returns **A** and **C** in a plain list, whereas `summary.dfm` returns all system matrices and additional residual and goodness of fit statistics—with a print method allowing full or compact printout.

Usage

```

## S3 method for class 'dfm'
print(x, digits = 4L, ...)

## S3 method for class 'dfm'
coef(object, ...)

## S3 method for class 'dfm'
logLik(object, ...)

## S3 method for class 'dfm'
summary(object, method = switch(object$em.method, none = "2s", "qml"), ...)

## S3 method for class 'dfm_summary'
print(x, digits = 4L, compact = sum(x$info["n"] > 15, x$info["n"] > 40), ...)

```

Arguments

<code>x</code> , object	an object class 'dfm'.
<code>digits</code>	integer. The number of digits to print out.
<code>...</code>	not used.
<code>method</code>	character. The factor estimates to use: one of "qml", "2s" or "pca".
<code>compact</code>	integer. Display a more compact printout: 0 prints everything, 1 omits the observation matrix C and residual covariance matrix <code>cov(resid(model))</code> , and 2 omits all disaggregated information on the input data. Sensible default are chosen for different sizes of the input dataset so as to limit large printouts.

Value

Summary information following a dynamic factor model estimation. `coef()` returns **A** and **C**.

See Also

[dfms-package](#)

Examples

```
mod <- DFM(diff(BM14_Q), 2, 3)
print(mod)
summary(mod)
```

tsnarmimp

Remove and Impute Missing Values in a Multivariate Time Series

Description

This function imputes missing values in a stationary multivariate time series using various methods, and removes cases with too many missing values.

Usage

```
tsnarmimp(
  X,
  max.missing = 0.8,
  na.rm.method = c("LE", "all"),
  na.impute = c("median.ma.spline", "median.ma", "median", "rnorm"),
  ma.terms = 3L
)
```

Arguments

<code>X</code>	a $T \times n$ numeric data matrix (incl. <code>ts</code> or <code>xts</code> objects) or data frame of stationary time series.
<code>max.missing</code>	numeric. Proportion of series missing for a case to be considered missing.
<code>na.rm.method</code>	character. Method to apply concerning missing cases selected through <code>max.missing</code> : "LE" only removes cases at the beginning or end of the sample, whereas "all" always removes missing cases.
<code>na.impute</code>	character. Method to impute missing values for the PCA estimates used to initialize the EM algorithm. Note that data are standardized (scaled and centered) beforehand. Available options are:
"median"	simple series-wise median imputation.
"rnorm"	imputation with random numbers drawn from a standard normal distribution.
"median.ma"	values are initially imputed with the median, but then a moving average is applied to smooth the e
"median.ma.spline"	"internal" missing values (not at the beginning or end of the sample) are imputed using a cubic spl
<code>ma.terms</code>	the order of the (2-sided) moving average applied in <code>na.impute</code> methods "median.ma" and "median.ma.spline".

Value

The imputed matrix `X_imp`, with attributes:

"missing"	a missingness matrix <code>W</code> matching the dimensions of <code>X_imp</code> .
"rm.rows"	and a vector of indices of rows (cases) with too many missing values that were removed.

See Also

[dfms-package](#)

Examples

```
library(xts)
str(tsnarmimp(BM14_M))
```

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