

Package ‘EstemPMM’

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Type Package

Title Polynomial Maximization Method for Non-Gaussian Regression

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Description Implements the Polynomial Maximization Method ('PMM') for parameter estimation in linear and time series models when error distributions deviate from normality. The 'PMM2' variant achieves lower variance parameter estimates compared to ordinary least squares ('OLS') when errors exhibit significant skewness. The 'PMM3' variant (S=3) targets symmetric platykurtic error distributions, reducing variance when excess kurtosis is negative. Includes automatic method selection ('pmm_dispatch'), linear regression, 'AR'/'MA'/'ARMA'/'ARIMA' models, and bootstrap inference. Methodology described in Zabolotnii, Warsza, and Tkachenko (2018) <[doi:10.1007/978-3-319-77179-3_75](https://doi.org/10.1007/978-3-319-77179-3_75)>, Zabolotnii, Tkachenko, and Warsza (2022) <[doi:10.1007/978-3-031-03502-9_37](https://doi.org/10.1007/978-3-031-03502-9_37)>, and Zabolotnii, Tkachenko, and Warsza (2023) <[doi:10.1007/978-3-031-25844-2_21](https://doi.org/10.1007/978-3-031-25844-2_21)>.

License GPL-3

Encoding UTF-8

Depends R (>= 3.5.0)

Imports methods, stats, graphics, utils

Suggests dplyr, ggplot2, gridExtra, testthat (>= 3.0.0), rmarkdown, knitr, MASS, numDeriv

LazyData true

RoxygenNote 7.3.3

Config/testthat/edition 3

URL <https://github.com/SZabolotnii/EstemPMM>

BugReports <https://github.com/SZabolotnii/EstemPMM/issues>

VignetteBuilder knitr

NeedsCompilation no

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AIC,PMM2fit-method *Calculate AIC for PMM2fit object*

Description

Calculate AIC for PMM2fit object

Usage

```
## S4 method for signature 'PMM2fit'
AIC(object, ..., k = 2)
```

Arguments

| | |
|--------|--|
| object | PMM2fit object |
| ... | Additional arguments (not used) |
| k | Penalty per parameter to be used; default is 2 |

Value

AIC value

AIC,PMM3fit-method *Calculate AIC for PMM3fit object*

Description

Calculate AIC for PMM3fit object

Usage

```
## S4 method for signature 'PMM3fit'
AIC(object, ..., k = 2)
```

Arguments

| | |
|--------|-----------------------------------|
| object | PMM3fit object |
| ... | Additional arguments (not used) |
| k | Penalty per parameter (default 2) |

Value

AIC value

| | |
|--------------------|--|
| AIC, TS3fit-method | <i>Calculate AIC for TS3fit object</i> |
|--------------------|--|

Description

Calculate AIC for TS3fit object

Usage

```
## S4 method for signature 'TS3fit'
AIC(object, ..., k = 2)
```

Arguments

| | |
|--------|-----------------------------------|
| object | TS3fit object |
| ... | Additional arguments (not used) |
| k | Penalty per parameter (default 2) |

Value

AIC value

| | |
|-----------------|--|
| ARIMAPMM2-class | <i>S4 class for storing PMM2 ARIMA model results</i> |
|-----------------|--|

Description

S4 class for storing PMM2 ARIMA model results

| | |
|-----------------|--|
| ARIMAPMM3-class | <i>S4 class for PMM3 ARIMA model results</i> |
|-----------------|--|

Description

S4 class for PMM3 ARIMA model results

arima_pmm2

Fit an ARIMA model using PMM2 (wrapper)

Description

Estimates autoregressive integrated moving-average model parameters using PMM2. ARIMA models extend ARMA to non-stationary series via differencing. PMM2 provides robust parameter estimates for the stationary ARMA component after differencing is applied.

Usage

```
arima_pmm2(
  x,
  order = c(1, 1, 1),
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification: - For AR models: a single integer (AR order) - For MA models: a single integer (MA order) - For ARMA models: vector c(p, q) (AR and MA orders) - For ARIMA models: vector c(p, d, q) (AR, differencing, and MA orders) |
| method | String: estimation method, one of "pmm2" (default), "css", "ml", "yw", "ols" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson), or "linearized" (specialized for MA/SMA models). |
| max_iter | Integer: maximum number of iterations for the algorithm |
| tol | Numeric: tolerance for convergence |
| include.mean | Logical: whether to include a mean (intercept) term |
| initial | List or vector of initial parameter estimates (optional) |
| na.action | Function for handling missing values, default is na.fail |
| regularize | Logical, add small values to diagonal for numerical stability |
| reg_lambda | Regularization parameter (if regularize=TRUE) |
| verbose | Logical: whether to print progress information |

Details

PMM2 Variants:

- "unified_global" (default): One-step correction from MLE/CSS estimates. Fast and reliable for most ARIMA specifications.
- "unified_iterative": Full Newton-Raphson optimization. Recommended for ARIMA models with complex dynamics or when unified_global shows residual non-Gaussianity.
- "linearized": First-order approximation. Not typically recommended for ARIMA unless MA component dominates.

Variant Selection Guidelines:

- For standard ARIMA(p,d,q): Use "unified_global" (default)
- For models with $d \geq 2$ or high orders: Try "unified_iterative"
- If MA component is large relative to AR: Consider "unified_iterative"

ARIMA Estimation Workflow:

1. Apply differencing of order d to achieve stationarity
2. Estimate ARMA(p,q) model on differenced series using PMM2
3. Return coefficients and diagnostics for the integrated model

Differencing Notes: The d parameter determines how many times the series is differenced. $d=0$ reduces to ARMA, $d=1$ handles unit root processes, $d=2$ is rare but useful for some economic series with trend acceleration.

Value

An S4 object of class ARIMAPMM2 containing fitted AR and MA coefficients, residual series, central moments, differencing order, intercept, original series, and convergence diagnostics.

See Also

[arma_pmm2](#), [sarima_pmm2](#), [ar_pmm2](#)

Examples

```
# Fit ARIMA(1,1,1) model to non-stationary series
x <- cumsum(arima.sim(n = 200, list(ar = 0.6, ma = -0.4)))
fit1 <- arima_pmm2(x, order = c(1, 1, 1))
coef(fit1)

# ARIMA(2,1,0) - random walk with AR(2) innovations
x2 <- cumsum(arima.sim(n = 250, list(ar = c(0.7, -0.3))))
fit2 <- arima_pmm2(x2, order = c(2, 1, 0), pmm2_variant = "unified_global")

# ARIMA(0,2,2) - double differencing with MA(2)
x3 <- cumsum(cumsum(arima.sim(n = 300, list(ma = c(0.5, 0.3))))))
fit3 <- arima_pmm2(x3, order = c(0, 2, 2), pmm2_variant = "unified_iterative")
```

`arima_pmm3`*Fit an ARIMA model using PMM3*

Description

Estimates ARIMA model parameters using PMM3.

Usage

```
arima_pmm3(  
  x,  
  order = c(1, 1, 1),  
  max_iter = 100,  
  tol = 1e-06,  
  adaptive = FALSE,  
  step_max = 5,  
  include.mean = TRUE,  
  initial = NULL,  
  na.action = na.fail,  
  verbose = FALSE  
)
```

Arguments

| | |
|---------------------------|---|
| <code>x</code> | Numeric vector of time series data |
| <code>order</code> | Numeric vector $c(p, d, q)$: AR, differencing, and MA orders |
| <code>max_iter</code> | Integer: maximum NR iterations (default 100) |
| <code>tol</code> | Numeric: convergence tolerance (default 1e-6) |
| <code>adaptive</code> | Logical: re-estimate kappa each iteration (default FALSE) |
| <code>step_max</code> | Numeric: maximum NR step size (default 5.0) |
| <code>include.mean</code> | Logical: include mean term (default TRUE) |
| <code>initial</code> | Optional initial parameter estimates |
| <code>na.action</code> | Function for handling missing values (default na.fail) |
| <code>verbose</code> | Logical: print progress information (default FALSE) |

Value

An S4 object of class ARIMAPMM3

See Also

[arima_pmm2](#), [arma_pmm3](#), [lm_pmm3](#)

Examples

```

set.seed(42)
x <- cumsum(arima.sim(n = 200, list(ar = 0.6),
                        innov = runif(200, -sqrt(3), sqrt(3))))
fit <- arima_pmm3(x, order = c(1, 1, 0))
coef(fit)

```

| | |
|----------------|---|
| ARMAPMM2-class | <i>S4 class for storing PMM2 ARMA model results</i> |
|----------------|---|

Description

S4 class for storing PMM2 ARMA model results

| | |
|----------------|---|
| ARMAPMM3-class | <i>S4 class for PMM3 ARMA model results</i> |
|----------------|---|

Description

S4 class for PMM3 ARMA model results

| | |
|-----------|---|
| arma_pmm2 | <i>Fit an ARMA model using PMM2 (wrapper)</i> |
|-----------|---|

Description

Estimates autoregressive moving-average model parameters using PMM2. ARMA models combine AR and MA components, capturing both direct past value dependencies and innovation structure. PMM2 leverages higher moments to improve parameter estimation accuracy.

Usage

```

arma_pmm2(
  x,
  order = c(1, 1),
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  initial = NULL,

```

```

na.action = na.fail,
regularize = TRUE,
reg_lambda = 1e-08,
verbose = FALSE
)

```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification: - For AR models: a single integer (AR order) - For MA models: a single integer (MA order) - For ARMA models: vector c(p, q) (AR and MA orders) - For ARIMA models: vector c(p, d, q) (AR, differencing, and MA orders) |
| method | String: estimation method, one of "pmm2" (default), "css", "ml", "yw", "ols" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson), or "linearized" (specialized for MA/SMA models). |
| max_iter | Integer: maximum number of iterations for the algorithm |
| tol | Numeric: tolerance for convergence |
| include.mean | Logical: whether to include a mean (intercept) term |
| initial | List or vector of initial parameter estimates (optional) |
| na.action | Function for handling missing values, default is na.fail |
| regularize | Logical, add small values to diagonal for numerical stability |
| reg_lambda | Regularization parameter (if regularize=TRUE) |
| verbose | Logical: whether to print progress information |

Details

PMM2 Variants:

- "unified_global" (default): One-step correction from MLE/CSS estimates. Balances speed and accuracy for ARMA models. Recommended for most use cases.
- "unified_iterative": Full Newton-Raphson optimization. Best for models with complex dynamics or strong non-Gaussian features.
- "linearized": First-order approximation. Generally not recommended for ARMA; better suited for pure MA/SMA models.

Variant Selection Guidelines:

- For ARMA(p,q) with $p, q \leq 2$: Use "unified_global" (default)
- For higher-order ARMA or ill-conditioned models: Try "unified_iterative"
- For quick exploration: Start with "unified_global"

ARMA Estimation Challenges: ARMA models have more parameters than pure AR or MA models, making estimation more sensitive to initialization and numerical stability. PMM2 benefits from robust moment-based constraints that help regularize the parameter space.

Value

An S4 object of class ARMAPMM2 containing fitted AR and MA coefficients, residuals, central moments, model specification, intercept, original series, and convergence diagnostics.

See Also

[ar_pmm2](#), [ma_pmm2](#), [arima_pmm2](#)

Examples

```
# Fit ARMA(2,1) model
x <- arima.sim(n = 250, list(ar = c(0.7, -0.3), ma = 0.5))
fit1 <- arma_pmm2(x, order = c(2, 1))
coef(fit1)

# Try iterative variant for better accuracy
fit2 <- arma_pmm2(x, order = c(2, 1), pmm2_variant = "unified_iterative")

# Higher-order ARMA
x2 <- arima.sim(n = 300, list(ar = c(0.6, -0.2), ma = c(0.4, 0.3)))
fit3 <- arma_pmm2(x2, order = c(2, 2), pmm2_variant = "unified_global")
```

arma_pmm3

Fit an ARMA model using PMM3

Description

Estimates ARMA model parameters using PMM3.

Usage

```
arma_pmm3(  
  x,  
  order = c(1, 1),  
  max_iter = 100,  
  tol = 1e-06,  
  adaptive = FALSE,  
  step_max = 5,  
  include.mean = TRUE,  
  initial = NULL,  
  na.action = na.fail,  
  verbose = FALSE  
)
```

Arguments

| | |
|---------------------------|--|
| <code>x</code> | Numeric vector of time series data |
| <code>order</code> | Numeric vector $c(p, q)$: AR and MA orders |
| <code>max_iter</code> | Integer: maximum NR iterations (default 100) |
| <code>tol</code> | Numeric: convergence tolerance (default $1e-6$) |
| <code>adaptive</code> | Logical: re-estimate kappa each iteration (default FALSE) |
| <code>step_max</code> | Numeric: maximum NR step size (default 5.0) |
| <code>include.mean</code> | Logical: include mean term (default TRUE) |
| <code>initial</code> | Optional initial parameter estimates |
| <code>na.action</code> | Function for handling missing values (default <code>na.fail</code>) |
| <code>verbose</code> | Logical: print progress information (default FALSE) |

Value

An S4 object of class ARMAPMM3

See Also

[arma_pmm2](#), [arima_pmm3](#), [lm_pmm3](#)

Examples

```
set.seed(42)
x <- arima.sim(n = 250, list(ar = 0.7, ma = -0.3),
              innov = runif(250, -sqrt(3), sqrt(3)))
fit <- arma_pmm3(x, order = c(1, 1))
coef(fit)
```

ARPM2-class

S4 class for storing PMM2 AR model results

Description

S4 class for storing PMM2 AR model results

ARPM3-class

S4 class for PMM3 AR model results

Description

S4 class for PMM3 AR model results

ar_pmm2

*Fit an AR model using PMM2 (wrapper)***Description**

Estimates autoregressive model parameters using the Pearson Moment Method (PMM2). PMM2 exploits third and fourth moment information to achieve more accurate parameter estimates than classical maximum likelihood, particularly for non-Gaussian innovations.

Usage

```
ar_pmm2(
  x,
  order = 1,
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification: - For AR models: a single integer (AR order) - For MA models: a single integer (MA order) - For ARMA models: vector c(p, q) (AR and MA orders) - For ARIMA models: vector c(p, d, q) (AR, differencing, and MA orders) |
| method | String: estimation method, one of "pmm2" (default), "css", "ml", "yw", "ols" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson), or "linearized" (specialized for MA/SMA models). |
| max_iter | Integer: maximum number of iterations for the algorithm |
| tol | Numeric: tolerance for convergence |
| include.mean | Logical: whether to include a mean (intercept) term |
| initial | List or vector of initial parameter estimates (optional) |
| na.action | Function for handling missing values, default is na.fail |
| regularize | Logical, add small values to diagonal for numerical stability |
| reg_lambda | Regularization parameter (if regularize=TRUE) |
| verbose | Logical: whether to print progress information |

Details**PMM2 Variants:**

- "unified_global" (default): One-step correction from MLE/CSS estimates. Fast and stable. Recommended for most AR models. Typical improvement: 3-5\
- "unified_iterative": Full Newton-Raphson optimization starting from classical estimates. More accurate but computationally intensive. Best for complex models with strong non-Gaussian features.
- "linearized": Uses first-order Taylor expansion. Not recommended for AR models; designed for MA/SMA where Jacobian computation is complex.

Variant Selection Guidelines:

- For AR(p) models: Use "unified_global" (default)
- If convergence issues occur: Try "unified_iterative"
- For highly skewed/heavy-tailed innovations: Use "unified_iterative"

Computational Characteristics:

- unified_global: ~2x slower than MLE (single correction step)
- unified_iterative: 5-10x slower than MLE (iterative refinement)
- linearized: ~1.5x slower than MLE (approximation-based)

Value

An S4 object of class ARPMM2 containing fitted autoregressive coefficients, residuals, central moment estimates (m2-m4), model order, intercept, original series, and convergence diagnostics.

References

Monte Carlo validation (R=50, n=200): Unified Iterative showed 2.9\ improvement over MLE for AR(1) models. See NEWS.md (Version 0.2.0) for details.

See Also

[ma_pmm2](#), [arma_pmm2](#), [arima_pmm2](#)

Examples

```
# Fit AR(2) model with default variant
x <- arima.sim(n = 200, list(ar = c(0.7, -0.3)))
fit1 <- ar_pmm2(x, order = 2)
coef(fit1)

# Compare variants
fit2 <- ar_pmm2(x, order = 2, pmm2_variant = "unified_iterative")
fit3 <- ar_pmm2(x, order = 2, pmm2_variant = "linearized")
```

`ar_pmm3`*Fit an AR model using PMM3*

Description

Estimates autoregressive model parameters using the Polynomial Maximization Method of order 3 (PMM3). Designed for symmetric platykurtic innovations.

Usage

```
ar_pmm3(  
  x,  
  order = 1,  
  max_iter = 100,  
  tol = 1e-06,  
  adaptive = FALSE,  
  step_max = 5,  
  include.mean = TRUE,  
  initial = NULL,  
  na.action = na.fail,  
  verbose = FALSE  
)
```

Arguments

| | |
|---------------------------|---|
| <code>x</code> | Numeric vector of time series data |
| <code>order</code> | Integer: AR order (default 1) |
| <code>max_iter</code> | Integer: maximum NR iterations (default 100) |
| <code>tol</code> | Numeric: convergence tolerance (default 1e-6) |
| <code>adaptive</code> | Logical: re-estimate kappa each iteration (default FALSE) |
| <code>step_max</code> | Numeric: maximum NR step size (default 5.0) |
| <code>include.mean</code> | Logical: include mean term (default TRUE) |
| <code>initial</code> | Optional initial parameter estimates |
| <code>na.action</code> | Function for handling missing values (default na.fail) |
| <code>verbose</code> | Logical: print progress information (default FALSE) |

Value

An S4 object of class ARPMM3

See Also

[ar_pmm2](#), [ma_pmm3](#), [lm_pmm3](#)

Examples

```
set.seed(42)
x <- arima.sim(n = 200, list(ar = 0.7),
               innov = runif(200, -sqrt(3), sqrt(3)))
fit <- ar_pmm3(x, order = 1)
coef(fit)
```

auto_mpg

Auto MPG Dataset

Description

Fuel consumption and vehicle characteristics for 398 automobiles from the 1970s and 1980s. This dataset is used in published PMM research to demonstrate both PMM2 (asymmetric residuals: MPG vs Weight) and PMM3 (symmetric platykurtic residuals: MPG vs Horsepower).

Usage

auto_mpg

Format

A data frame with 398 rows and 9 variables:

mpg Miles per gallon (fuel efficiency)
cylinders Number of cylinders (4, 6, or 8)
displacement Engine displacement (cubic inches)
horsepower Engine horsepower (6 missing values)
weight Vehicle weight (pounds)
acceleration Time to accelerate from 0 to 60 mph (seconds)
model_year Model year (70-82, i.e., 1970-1982)
origin Origin (1 = American, 2 = European, 3 = Japanese)
car_name Car model name

Details

Three regression examples from published PMM papers:

- **MPG vs Acceleration** (PMM2, linear): residuals have $\gamma_3 = 0.49$, $g_2 = 0.86$ (Zabolotnii et al., 2018)
- **MPG vs Weight** (PMM2, quadratic): residuals have $\gamma_3 = 0.8$, $g_2 = 0.83$ (Zabolotnii et al., 2025)
- **MPG vs Horsepower** (PMM3, quadratic): residuals have $\gamma_3 \sim 0.2$, $\gamma_4 = 1.3$, $g_3 = 0.89$ (Zabolotnii et al., 2025)

Source

UCI Machine Learning Repository <https://archive.ics.uci.edu/dataset/9/auto+mpg>

References

Quinlan, J.R. (1993). Combining Instance-Based and Model-Based Learning. In Proceedings on the Tenth International Conference of Machine Learning, 236-243.

Zabolotnii S., Warsza Z.L., Tkachenko O. (2018) Polynomial Estimation of Linear Regression Parameters for the Asymmetric PDF of Errors. Springer AISC, vol 743. doi:10.1007/978331977179-3_75

Examples

```
data(auto_mpg)
# PMM2 example: MPG vs Acceleration (asymmetric residuals)
fit_ols <- lm(mpg ~ acceleration, data = auto_mpg)
pmm_skewness(residuals(fit_ols)) # gamma3 ~ 0.5 -> PMM2
pmm_dispatch(residuals(fit_ols))
fit_pmm2 <- lm_pmm2(mpg ~ acceleration, data = auto_mpg, na.action = na.omit)
coef(fit_pmm2) # compare with coef(fit_ols)
```

BasePMM2-class

Base S4 class for storing PMM2 model results

Description

Base S4 class for storing PMM2 model results

Slots

`coefficients` numeric vector of estimated parameters

`residuals` numeric vector of final residuals

`m2` numeric second central moment of initial residuals

`m3` numeric third central moment of initial residuals

`m4` numeric fourth central moment of initial residuals

`convergence` logical or integer code indicating whether algorithm converged

`iterations` numeric number of iterations performed

`call` original function call

coef,PMM2fit-method *Extract coefficients from PMM2fit object*

Description

Extract coefficients from PMM2fit object

Usage

```
## S4 method for signature 'PMM2fit'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM2fit object |
| ... | Additional arguments (not used) |

Value

Vector of coefficients

coef,PMM3fit-method *Extract coefficients from PMM3fit object*

Description

Extract coefficients from PMM3fit object

Usage

```
## S4 method for signature 'PMM3fit'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM3fit object |
| ... | Additional arguments (not used) |

Value

Vector of coefficients

coef, SARPMM2-method *Extract coefficients from SARPMM2 object*

Description

Extract coefficients from SARPMM2 object

Usage

```
## S4 method for signature 'SARPMM2'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | SARPMM2 object |
| ... | Additional arguments (not used) |

Value

Named numeric vector of coefficients

coef, SMAPMM2-method *Extract coefficients from SMAPMM2 object*

Description

Extract coefficients from SMAPMM2 object

Usage

```
## S4 method for signature 'SMAPMM2'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | SMAPMM2 object |
| ... | Additional arguments (not used) |

Value

Named vector of seasonal MA coefficients

coef,TS2fit-method *Extract coefficients from TS2fit object*

Description

Extract coefficients from TS2fit object

Usage

```
## S4 method for signature 'TS2fit'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS2fit object |
| ... | Additional arguments (not used) |

Value

Named vector of coefficients

coef,TS3fit-method *Extract coefficients from TS3fit object*

Description

Extract coefficients from TS3fit object

Usage

```
## S4 method for signature 'TS3fit'  
coef(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS3fit object |
| ... | Additional arguments (not used) |

Value

Named vector of coefficients

compare_arma_methods *Compare ARMA methods*

Description

Compare ARMA methods

Usage

```
compare_arma_methods(  
  x,  
  order = c(1, 1, 1),  
  include.mean = TRUE,  
  pmm2_args = list()  
)
```

Arguments

| | |
|--------------|--|
| x | Numeric vector of time series data |
| order | Model order specification (see ts_pmm2 for format) |
| include.mean | Logical, whether to include intercept term |
| pmm2_args | List of additional arguments to pass to ts_pmm2() |

Value

A named list containing the fitted objects for each estimation approach (e.g., YW/OLS/MLE or CSS/ML alongside PMM2) plus two data frames: `coefficients` (side-by-side parameter estimates) and `residual_stats` (residual RSS, MAE, skewness, and kurtosis).

compare_arma_methods *Compare ARMA methods*

Description

Compare ARMA methods

Usage

```
compare_arma_methods(  
  x,  
  order = c(1, 1),  
  include.mean = TRUE,  
  pmm2_args = list()  
)
```

Arguments

| | |
|--------------|--|
| x | Numeric vector of time series data |
| order | Model order specification (see ts_pmm2 for format) |
| include.mean | Logical, whether to include intercept term |
| pmm2_args | List of additional arguments to pass to ts_pmm2() |

Value

A named list containing the fitted objects for each estimation approach (e.g., YW/OLS/MLE or CSS/ML alongside PMM2) plus two data frames: `coefficients` (side-by-side parameter estimates) and `residual_stats` (residual RSS, MAE, skewness, and kurtosis).

| | |
|--------------------|---------------------------|
| compare_ar_methods | <i>Compare AR methods</i> |
|--------------------|---------------------------|

Description

Compare AR methods

Usage

```
compare_ar_methods(x, order = 1, include.mean = TRUE, pmm2_args = list())
```

Arguments

| | |
|--------------|--|
| x | Numeric vector of time series data |
| order | Model order specification (see ts_pmm2 for format) |
| include.mean | Logical, whether to include intercept term |
| pmm2_args | List of additional arguments to pass to ts_pmm2() |

Value

A named list containing the fitted objects for each estimation approach (e.g., YW/OLS/MLE or CSS/ML alongside PMM2) plus two data frames: `coefficients` (side-by-side parameter estimates) and `residual_stats` (residual RSS, MAE, skewness, and kurtosis).

| | |
|--------------------|---------------------------|
| compare_ma_methods | <i>Compare MA methods</i> |
|--------------------|---------------------------|

Description

Compare MA methods

Usage

```
compare_ma_methods(x, order = 1, include.mean = TRUE, pmm2_args = list())
```

Arguments

| | |
|--------------|--|
| x | Numeric vector of time series data |
| order | Model order specification (see ts_pmm2 for format) |
| include.mean | Logical, whether to include intercept term |
| pmm2_args | List of additional arguments to pass to ts_pmm2() |

Value

A named list containing the fitted objects for each estimation approach (e.g., YW/OLS/MLE or CSS/ML alongside PMM2) plus two data frames: `coefficients` (side-by-side parameter estimates) and `residual_stats` (residual RSS, MAE, skewness, and kurtosis).

| | |
|---------------------|---|
| compare_sar_methods | <i>Compare SAR model estimation methods</i> |
|---------------------|---|

Description

Compares different estimation methods (OLS, PMM2, CSS, ML) for SAR models on the same data.

Usage

```
compare_sar_methods(
  x,
  order = c(1, 1),
  period = 12,
  methods = c("ols", "pmm2", "css"),
  verbose = TRUE
)
```

Arguments

| | |
|---------|---|
| x | Time series data |
| order | Model order $c(p, P)$ for SAR specification |
| period | Seasonal period |
| methods | Character vector of methods to compare (default: <code>c("ols", "pmm2", "css")</code>) |
| verbose | Logical: print results to console (default TRUE) |

Value

Data frame with comparison results (invisibly)

Examples

```
set.seed(42)
y <- arima.sim(n = 120,
  model = list(order = c(1, 0, 0), ar = 0.7,
    seasonal = list(order = c(1, 0, 0), ar = 0.5, period = 12)))
compare_sar_methods(y, order = c(1, 1), period = 12)
```

| | |
|--------------------|---|
| compare_ts_methods | <i>Compare PMM2 with classical time series estimation methods</i> |
|--------------------|---|

Description

Compare PMM2 with classical time series estimation methods

Usage

```
compare_ts_methods(
  x,
  order,
  model_type = c("ar", "ma", "arma", "arima"),
  include.mean = TRUE,
  pmm2_args = list()
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification (see <code>ts_pmm2</code> for format) |
| model_type | Model type: "ar", "ma", "arma", or "arima" |
| include.mean | Logical, whether to include intercept term |
| pmm2_args | List of additional arguments to pass to <code>ts_pmm2()</code> |

Value

A named list containing the fitted objects for each estimation approach (e.g., YW/OLS/MLE or CSS/ML alongside PMM2) plus two data frames: `coefficients` (side-by-side parameter estimates) and `residual_stats` (residual RSS, MAE, skewness, and kurtosis).

| | |
|------------------|------------------------------|
| compare_with_ols | <i>Compare PMM2 with OLS</i> |
|------------------|------------------------------|

Description

Compare PMM2 with OLS

Usage

```
compare_with_ols(formula, data, pmm2_args = list())
```

Arguments

| | |
|-----------|---|
| formula | Model formula |
| data | Data frame |
| pmm2_args | List of arguments to pass to <code>lm_pmm2()</code> |

Value

List with OLS and PMM2 fit objects

Examples

```
result <- compare_with_ols(mpg ~ wt, data = mtcars)
```

| | |
|-----------------|--|
| compute_moments | <i>Calculate moments and cumulants of error distribution</i> |
|-----------------|--|

Description

Calculate moments and cumulants of error distribution

Usage

```
compute_moments(errors)
```

Arguments

| | |
|--------|--------------------------|
| errors | numeric vector of errors |
|--------|--------------------------|

Value

list with moments, cumulants and theoretical variance reduction coefficient

compute_moments_pmm3 *Compute central moments for PMM3 from residuals*

Description

Computes the second, fourth, and sixth central moments, along with standardised cumulant coefficients gamma3, gamma4, gamma6, the theoretical efficiency factor g3, and the moment ratio kappa used in the PMM3 solver.

Usage

```
compute_moments_pmm3(residuals)
```

Arguments

residuals numeric vector of residuals (typically from OLS)

Value

A list with components:

| | |
|-----------------|--|
| m2 | Second central moment |
| m4 | Fourth central moment |
| m6 | Sixth central moment |
| gamma3 | Skewness coefficient (for symmetry check) |
| gamma4 | Excess kurtosis |
| gamma6 | Sixth-order cumulant coefficient |
| g3 | Theoretical variance reduction factor |
| kappa | Moment ratio for NR solver (NA if near-Gaussian) |
| improvement_pct | Expected variance reduction percentage |

`compute_pmm2_components`*Compute PMM2 weights and components*

Description

Compute PMM2 weights and components

Usage

```
compute_pmm2_components(residuals)
```

Arguments

`residuals` Residual vector

Value

List with moments and PMM2 parameters (`gamma3`, `gamma4`, `weights`)

`create_sarma_matrix`*Create design matrix for seasonal ARMA model*

Description

Constructs a design matrix for Seasonal ARMA (SARMA) models, combining non-seasonal and seasonal AR and MA components.

Usage

```
create_sarma_matrix(  
  x,  
  residuals,  
  p = 0,  
  P = 0,  
  q = 0,  
  Q = 0,  
  s = 12,  
  multiplicative = FALSE  
)
```

Arguments

| | |
|----------------|---|
| x | Numeric vector (centered time series) |
| residuals | Numeric vector (initial residuals/innovations) |
| p | Non-seasonal AR order (default 0) |
| P | Seasonal AR order (default 0) |
| q | Non-seasonal MA order (default 0) |
| Q | Seasonal MA order (default 0) |
| s | Seasonal period (e.g., 12 for monthly data) |
| multiplicative | Logical, include multiplicative cross-terms (default FALSE) |

Details

For a SARMA(p,q) x (P,Q)_s model:

$$y_t = \phi_1 y_{t-1} + \dots + \phi_p y_{t-p} + \Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps} + \theta_1 \epsilon_{t-1} + \dots + \theta_q \epsilon_{t-q} + \Theta_1 \epsilon_{t-s} + \dots + \Theta_Q \epsilon_{t-Qs} + \epsilon_t.$$

Where:

- p, P are non-seasonal and seasonal AR orders
- q, Q are non-seasonal and seasonal MA orders
- s is the seasonal period

Value

Design matrix with lagged values. Columns are ordered as:

- First p columns: non-seasonal AR lags (y_{t-1}, \dots, y_{t-p})
- Next P columns: seasonal AR lags (y_{t-s}, \dots, y_{t-Ps})
- Next q columns: non-seasonal MA lags ($\epsilon_{t-1}, \dots, \epsilon_{t-q}$)
- Next Q columns: seasonal MA lags ($\epsilon_{t-s}, \dots, \epsilon_{t-Qs}$)
- If multiplicative=TRUE: AR cross-terms then MA cross-terms

Examples

```
# Simple SARMA(1,0) x (1,0)_12 model (AR+SAR, no MA)
x <- rnorm(120)
residuals <- rnorm(120)
X <- create_sarma_matrix(x, residuals, p = 1, P = 1, q = 0, Q = 0, s = 12)

# Full SARMA(1,1) x (1,1)_12 model
X <- create_sarma_matrix(x, residuals, p = 1, P = 1, q = 1, Q = 1, s = 12)
```

create_sar_matrix *Create design matrix for seasonal AR model*

Description

Constructs a design matrix for Seasonal Autoregressive (SAR) models, optionally including non-seasonal AR lags and multiplicative cross-terms.

Usage

```
create_sar_matrix(x, p = 0, P = 1, s = 12, multiplicative = FALSE)
```

Arguments

`x` Numeric vector (centered time series)

`p` Non-seasonal AR order (default 0)

`P` Seasonal AR order (must be positive)

`s` Seasonal period (e.g., 12 for monthly data)

`multiplicative` Logical, include multiplicative cross-terms (default FALSE)

Details

For a SAR(P)_s model:

$$y_t = \Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps} + \epsilon_t.$$

For an additive AR(p)+SAR(P)_s model:

$$y_t = \phi_1 y_{t-1} + \dots + \phi_p y_{t-p} + \Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps} + \epsilon_t.$$

For a multiplicative AR(p) x SAR(P)_s model (`multiplicative = TRUE`): Includes cross-terms such as y_{t-1-s} , y_{t-1-2s} , etc.

Value

Design matrix with lagged values. Columns are:

- First p columns: non-seasonal lags (y_{t-1}, \dots, y_{t-p})
- Next P columns: seasonal lags (y_{t-s}, \dots, y_{t-Ps})
- If `multiplicative = TRUE`: additional $p \times P$ columns for cross-terms

Examples

```
# Simple SAR(1)_12 model
x <- rnorm(120)
X <- create_sar_matrix(x, p = 0, P = 1, s = 12)

# AR(1) + SAR(1)_12 additive model
X <- create_sar_matrix(x, p = 1, P = 1, s = 12)

# AR(1) x SAR(1)_12 multiplicative model
X <- create_sar_matrix(x, p = 1, P = 1, s = 12, multiplicative = TRUE)
```

DCOILWTICO

WTI Crude Oil Prices

Description

Daily spot prices for West Texas Intermediate (WTI) crude oil in U.S. dollars per barrel.

Usage

```
DCOILWTICO
```

Format

A data frame with observations for each trading day:

observation_date Date of observation in YYYY-MM-DD format

DCOILWTICO Crude Oil Price: West Texas Intermediate (WTI) in USD per barrel

Source

Federal Reserve Economic Data (FRED), Federal Reserve Bank of St. Louis <https://fred.stlouisfed.org/series/DCOILWTICO>

Examples

```
data(DCOILWTICO)
head(DCOILWTICO)
summary(DCOILWTICO$DCOILWTICO)
```

`djia2002`*Dow Jones Industrial Average Daily Data (July-December 2002)*

Description

Daily closing prices and changes of the Dow Jones Industrial Average for the second half of 2002. Used in published PMM2 research to demonstrate AR(1) estimation with asymmetric innovations.

Usage`djia2002`**Format**

A data frame with 127 rows and 3 variables:

date Trading date

close DJIA closing price (USD)

change Daily change in closing price (first difference; NA for the first observation)

Details

The daily changes exhibit positive skewness, making this dataset suitable for PMM2 estimation. In the original paper, an AR(1) model fitted to the change series yielded PMM2 coefficient $a_1 = -0.43$ versus OLS $a_1 = -0.49$, with $g_2 = 0.77$ (23\

Source

Yahoo Finance via the `quantmod` R package.

References

Zabolotnii S., Tkachenko O., Warsza Z.L. (2022) Application of the Polynomial Maximization Method for Estimation Parameters of Autoregressive Models with Asymmetric Innovations. Springer AISC, vol 1427. [doi:10.1007/9783031035029_37](https://doi.org/10.1007/9783031035029_37)

Examples

```
data(djia2002)
# AR(1) with PMM2
changes <- na.omit(djia2002$change)
pmm_skewness(changes) # positive skewness -> PMM2
fit <- ar_pmm2(changes, order = 1)
summary(fit)
```

fitted,PMM2fit-method *Extract fitted values from PMM2fit object*

Description

Extract fitted values from PMM2fit object

Usage

```
## S4 method for signature 'PMM2fit'
fitted(object, data = NULL, ...)
```

Arguments

| | |
|--------|--|
| object | PMM2fit object |
| data | Optional data source for model reconstruction, if object does not contain saved data |
| ... | Additional arguments (not used) |

Value

Vector of fitted values

fitted,PMM3fit-method *Extract fitted values from PMM3fit object*

Description

Extract fitted values from PMM3fit object

Usage

```
## S4 method for signature 'PMM3fit'
fitted(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM3fit object |
| ... | Additional arguments (not used) |

Value

Vector of fitted values

fitted,TS2fit-method *Extract fitted values from TS2fit object*

Description

Extract fitted values from TS2fit object

Usage

```
## S4 method for signature 'TS2fit'  
fitted(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS2fit object |
| ... | Additional arguments (not used) |

Value

Vector of fitted values

fitted,TS3fit-method *Extract fitted values from TS3fit object*

Description

Extract fitted values from TS3fit object

Usage

```
## S4 method for signature 'TS3fit'  
fitted(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS3fit object |
| ... | Additional arguments (not used) |

Value

Vector of fitted values

get_sarimax_jacobian *Calculate SARIMAX Jacobian (Numerical)*

Description

Calculate SARIMAX Jacobian (Numerical)

Usage

```
get_sarimax_jacobian(theta, ...)
```

Arguments

| | |
|-------|---|
| theta | Parameters |
| ... | Arguments passed to get_sarimax_residuals |

Value

Jacobian matrix (n x p)

get_sarimax_residuals *Calculate SARIMAX Residuals*

Description

Calculate SARIMAX Residuals

Usage

```
get_sarimax_residuals(
  theta,
  y,
  xreg = NULL,
  order = c(0, 0, 0),
  seasonal = list(order = c(0, 0, 0), period = NA),
  include.mean = TRUE
)
```

Arguments

| | |
|--------------|---|
| theta | Combined vector of parameters (AR, MA, SAR, SMA, Intercept, Regressors) |
| y | Time series data |
| xreg | Exogenous regressors (optional) |
| order | ARIMA order c(p, d, q) |
| seasonal | Seasonal order c(P, D, Q) |
| include.mean | Boolean, whether to include mean/intercept |
| period | Seasonal period |

Value

Vector of residuals

| | |
|---------|---|
| lm_pmm2 | <i>PMM2: Main function for PMM2 (S=2)</i> |
|---------|---|

Description

Fits a linear model using the Polynomial Maximization Method (order 2), which is robust to non-Gaussian errors.

Usage

```
lm_pmm2(
  formula,
  data,
  max_iter = 50,
  tol = 1e-06,
  regularize = TRUE,
  reg_lambda = 1e-08,
  na.action = na.fail,
  weights = NULL,
  verbose = FALSE
)
```

Arguments

| | |
|------------|--|
| formula | R formula for the model |
| data | data.frame containing variables in the formula |
| max_iter | integer: maximum number of iterations for the algorithm |
| tol | numeric: tolerance for convergence |
| regularize | logical: add small value to diagonal for numerical stability |
| reg_lambda | numeric: regularization parameter (if regularize=TRUE) |
| na.action | function for handling missing values, default is na.fail |
| weights | optional weight vector (not yet implemented) |
| verbose | logical: whether to print progress information |

Details

The PMM2 algorithm works as follows:

1. Fits ordinary least squares (OLS) regression to obtain initial estimates
2. Computes central moments (m2, m3, m4) from OLS residuals
3. Iteratively improves parameter estimates using a gradient-based approach

PMM2 is especially useful when error terms are not Gaussian.

Value

S4 object of class PMM2fit

Examples

```
set.seed(123)
n <- 80
x <- rnorm(n)
y <- 2 + 3 * x + rt(n, df = 3)
dat <- data.frame(y = y, x = x)

fit <- lm_pmm2(y ~ x, data = dat)
summary(fit, formula = y ~ x, data = dat)
```

| | |
|---------|--|
| lm_pmm3 | <i>PMM3: Fit linear model using Polynomial Maximization Method (S=3)</i> |
|---------|--|

Description

Fits a linear model using PMM3, which is designed for symmetric platykurtic error distributions. Uses a cubic stochastic polynomial with Newton-Raphson solver.

Usage

```
lm_pmm3(
  formula,
  data,
  max_iter = 100,
  tol = 1e-06,
  adaptive = FALSE,
  step_max = 5,
  na.action = na.fail,
  verbose = FALSE
)
```

Arguments

| | |
|-----------|---|
| formula | R formula for the model |
| data | data.frame containing variables in the formula |
| max_iter | integer: maximum number of NR iterations (default 100) |
| tol | numeric: convergence tolerance (default 1e-6) |
| adaptive | logical: re-estimate kappa each iteration (default FALSE) |
| step_max | numeric: maximum NR step size (default 5.0) |
| na.action | function for handling missing values (default na.fail) |
| verbose | logical: print progress information (default FALSE) |

Details

The PMM3 algorithm works as follows:

1. Fits OLS regression to obtain initial estimates
2. Computes central moments (m_2 , m_4 , m_6) from OLS residuals
3. Checks symmetry: warns if $|\text{gamma}_3| > 0.3$ (PMM2 may be more appropriate)
4. Computes moment ratio $\text{kappa} = (m_6 - 3m_4m_2) / (m_4 - 3m_2^2)$
5. Iterates Newton-Raphson with score $Z = X'(\text{eps} * (\text{kappa} - \text{eps}^2))$

PMM3 achieves lower variance than OLS when errors are symmetric and platykurtic ($\text{gamma}_4 < 0$), with theoretical efficiency $g_3 = 1 - \text{gamma}_4^2 / (6 + 9 * \text{gamma}_4 + \text{gamma}_6)$.

Value

S4 object of class PMM3fit

Examples

```
set.seed(123)
n <- 100
x <- rnorm(n)
y <- 2 + 3 * x + runif(n, -sqrt(3), sqrt(3))
dat <- data.frame(y = y, x = x)

fit <- lm_pmm3(y ~ x, data = dat)
summary(fit)
```

MAPMM2-class

S4 class for storing PMM2 MA model results

Description

S4 class for storing PMM2 MA model results

MAPMM3-class

S4 class for PMM3 MA model results

Description

S4 class for PMM3 MA model results

ma_pmm2

*Fit an MA model using PMM2 (wrapper)***Description**

Estimates moving-average model parameters using the Pearson Moment Method (PMM2). For MA models, PMM2 can achieve substantial improvements over MLE, particularly when innovations are non-Gaussian. Monte Carlo experiments showed up to 23\ reduction for MA(1) models.

Usage

```
ma_pmm2(
  x,
  order = 1,
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification: - For AR models: a single integer (AR order) - For MA models: a single integer (MA order) - For ARMA models: vector c(p, q) (AR and MA orders) - For ARIMA models: vector c(p, d, q) (AR, differencing, and MA orders) |
| method | String: estimation method, one of "pmm2" (default), "css", "ml", "yw", "ols" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson), or "linearized" (specialized for MA/SMA models, recommended for MA). |
| max_iter | Integer: maximum number of iterations for the algorithm |
| tol | Numeric: tolerance for convergence |
| include.mean | Logical: whether to include a mean (intercept) term |
| initial | List or vector of initial parameter estimates (optional) |
| na.action | Function for handling missing values, default is na.fail |
| regularize | Logical, add small values to diagonal for numerical stability |
| reg_lambda | Regularization parameter (if regularize=TRUE) |
| verbose | Logical: whether to print progress information |

Details

PMM2 Variants:

- "unified_global" (default): One-step correction from MLE/CSS estimates. Fast and reliable. Typical improvement: 15-23%
- "unified_iterative": Full Newton-Raphson optimization. Slower but can achieve better accuracy for complex MA models.
- "linearized": Uses first-order Taylor expansion around MLE. **Recommended for MA/SMA models** as it avoids complex Jacobian computation while maintaining accuracy. Fastest option for MA models.

Variant Selection Guidelines:

- For MA(q) models: Use "linearized" (fastest, MA-optimized)
- If you need maximum accuracy: Try "unified_global" (best MSE)
- For exploration: Compare all three variants

Computational Characteristics:

- linearized: ~1.5x slower than MLE (recommended)
- unified_global: ~2x slower than MLE (high accuracy)
- unified_iterative: 5-10x slower than MLE (iterative)

Why MA Models Benefit Most: MA parameter estimation from MLE has known numerical difficulties due to non-identifiability and flat likelihood regions. PMM2 uses moment constraints that better resolve these issues, leading to larger improvements than for AR models.

Value

An S4 object of class MAPMM2 containing moving-average coefficients, residual innovations, central moments, model order, intercept, original series, and convergence diagnostics.

References

Monte Carlo validation (R=50, n=200): Unified Global showed 23.0% improvement over MLE for MA(1) models - the largest improvement among all model types. See NEWS.md (Version 0.2.0) for full comparison.

See Also

[ar_pmm2](#), [arma_pmm2](#), [sma_pmm2](#)

Examples

```
# Fit MA(1) model with linearized variant (recommended)
x <- arima.sim(n = 200, list(ma = 0.6))
fit1 <- ma_pmm2(x, order = 1, pmm2_variant = "linearized")
coef(fit1)

# Compare with unified_global (best accuracy)
```

```
fit2 <- ma_pmm2(x, order = 1, pmm2_variant = "unified_global")

# Higher-order MA
x2 <- arima.sim(n = 300, list(ma = c(0.7, -0.4, 0.2)))
fit3 <- ma_pmm2(x2, order = 3, pmm2_variant = "linearized")
```

ma_pmm3

Fit an MA model using PMM3

Description

Estimates moving-average model parameters using PMM3.

Usage

```
ma_pmm3(
  x,
  order = 1,
  max_iter = 100,
  tol = 1e-06,
  adaptive = FALSE,
  step_max = 5,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Integer: AR order (default 1) |
| max_iter | Integer: maximum NR iterations (default 100) |
| tol | Numeric: convergence tolerance (default 1e-6) |
| adaptive | Logical: re-estimate kappa each iteration (default FALSE) |
| step_max | Numeric: maximum NR step size (default 5.0) |
| include.mean | Logical: include mean term (default TRUE) |
| initial | Optional initial parameter estimates |
| na.action | Function for handling missing values (default na.fail) |
| verbose | Logical: print progress information (default FALSE) |

Value

An S4 object of class MAPMM3

See Also

[ma_pmm2](#), [ar_pmm3](#), [lm_pmm3](#)

Examples

```
set.seed(42)
x <- arima.sim(n = 200, list(ma = 0.6),
               innov = runif(200, -sqrt(3), sqrt(3)))
fit <- ma_pmm3(x, order = 1)
coef(fit)
```

plot,PMM2fit,missing-method

Plot diagnostic plots for PMM2fit object

Description

Plot diagnostic plots for PMM2fit object

Usage

```
## S4 method for signature 'PMM2fit,missing'
plot(x, y, which = 1:4, ...)
```

Arguments

- x PMM2fit object
- y Not used (compatibility with generic)
- which Set of plots to display (values 1-4)
- ... Additional arguments passed to plotting functions

Value

Invisibly returns the input object

plot,PMM3fit,missing-method

Plot diagnostic plots for PMM3fit object

Description

Plot diagnostic plots for PMM3fit object

Usage

```
## S4 method for signature 'PMM3fit,missing'
plot(x, y, which = 1:4, ...)
```

Arguments

| | |
|-------|---|
| x | PMM3fit object |
| y | Not used (compatibility with generic) |
| which | Set of plots to display (values 1-4) |
| ... | Additional arguments passed to plotting functions |

Value

Invisibly returns the input object

plot,TS2fit,missing-method

Build diagnostic plots for TS2fit objects

Description

Build diagnostic plots for TS2fit objects

Usage

```
## S4 method for signature 'TS2fit,missing'
plot(x, y, which = c(1:4), ...)
```

Arguments

| | |
|-------|--|
| x | TS2fit object |
| y | Not used (for S4 method compatibility) |
| which | Integer vector indicating which plots to produce |
| ... | additional arguments passed to plot functions |

Value

Invisibly returns x

plot,TS3fit,missing-method
Plot diagnostic plots for TS3fit object

Description

Plot diagnostic plots for TS3fit object

Usage

```
## S4 method for signature 'TS3fit,missing'  
plot(x, y, which = 1:4, ...)
```

Arguments

| | |
|-------|--------------------------------------|
| x | TS3fit object |
| y | Not used |
| which | Set of plots to display (values 1-4) |
| ... | Additional arguments |

Value

Invisibly returns the input object

plot_pmm2_bootstrap *Plot bootstrap distributions for PMM2 fit*

Description

Plot bootstrap distributions for PMM2 fit

Usage

```
plot_pmm2_bootstrap(object, coefficients = NULL)
```

Arguments

| | |
|--------------|---|
| object | Result from pmm2_inference |
| coefficients | Which coefficients to plot, defaults to all |

Value

Invisibly returns histogram information

PMM2fit-class

S4 class for storing PMM2 regression model results

Description

Class for storing results of linear model estimation using PMM2

Slots

`coefficients` numeric vector of estimated parameters

`residuals` numeric vector of final residuals

`m2` numeric second central moment of initial residuals

`m3` numeric third central moment of initial residuals

`m4` numeric fourth central moment of initial residuals

`convergence` logical or integer code indicating whether algorithm converged

`iterations` numeric number of iterations performed

`call` original function call

Slots

coefficients Estimated coefficients

residuals Final residuals

m2 Second central moment

m3 Third central moment

m4 Fourth central moment

convergence Convergence status

iterations Number of iterations performed

call Original call

pmm2_inference

Bootstrap inference for PMM2 fit

Description

Bootstrap inference for PMM2 fit

Usage

```
pmm2_inference(
  object,
  formula,
  data,
  B = 200,
  seed = NULL,
  parallel = FALSE,
  cores = NULL
)
```

Arguments

| | |
|----------|--|
| object | object of class PMM2fit |
| formula | the same formula that was used initially |
| data | data frame that was used initially |
| B | number of bootstrap replications |
| seed | (optional) for reproducibility |
| parallel | logical, whether to use parallel computing |
| cores | number of cores to use for parallel computing, defaults to auto-detect |

Value

data.frame with columns: Estimate, Std.Error, t.value, p.value

pmm2_monte_carlo_compare

Monte Carlo comparison of PMM2 estimation methods

Description

Function generates time series for given models, repeatedly estimates parameters using different methods and compares their accuracy by MSE criterion. Additionally outputs theoretical and empirical characteristics of the innovation distribution (skewness, excess kurtosis, theoretical gain of PMM2).

Usage

```
pmm2_monte_carlo_compare(
  model_specs,
  methods = c("css", "pmm2"),
  n,
  n_sim,
  innovations = list(type = "gaussian"),
  seed = NULL,
```

```

include.mean = TRUE,
progress = interactive(),
verbose = FALSE
)

```

Arguments

| | |
|--------------|---|
| model_specs | List of model specifications. Each element must contain: model "ar", "ma" or "arma" order order (for AR/MA) or vector c(p, q) for ARMA theta numeric vector of true parameters; for ARMA a list list(ar = ..., ma = ...) label (optional) model name in report innovations (optional) description of innovation distribution: list(type = "gamma", shape = 2), list(type = "student_t", df = 5), etc. Can also pass an arbitrary generation function via generator. |
| methods | Vector of estimation methods (e.g., c("css", "pmm2")). The first method is considered baseline for relative MSE calculation. |
| n | Sample size for simulation. |
| n_sim | Number of Monte Carlo experiments. |
| innovations | Function or distribution description, used by default for all models (if not specified in spec). |
| seed | Initial seed for random number generator (optional). |
| include.mean | Logical flag: whether to include intercept during estimation. |
| progress | Logical flag: print Monte Carlo progress. |
| verbose | Whether to print diagnostic messages on failures. |

Value

List with three components:

- parameter_results** MSE and relative MSE for each parameter
- summary** Averaged MSE over parameters for each model/method
- gain** Comparison of theoretical and empirical PMM2 gain

pmm2_nonlinear_iterative

Universal PMM2 estimator (Iterative)

Description

Universal PMM2 estimator (Iterative)

Usage

```
pmm2_nonlinear_iterative(
  theta_init,
  fn_residuals,
  fn_jacobian = NULL,
  max_iter = 100,
  tol = 1e-06,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|--|
| theta_init | Initial parameter values |
| fn_residuals | Function(theta) returning the residual vector |
| fn_jacobian | Function(theta) returning the Jacobian matrix (n x p). $J_{ij} = d(\hat{y}_i)/d(\theta_j) = -d(\epsilon_i)/d(\theta_j)$. If NULL, numerical Jacobian via numDeriv is used |
| max_iter | Maximum number of iterations |
| tol | Convergence tolerance |
| verbose | Print progress |

Value

List with results (theta, residuals, convergence, etc.)

pmm2_nonlinear_onestep

Universal PMM2 estimator (One-step Global)

Description

Applies a single PMM2 correction to classical estimation results.

Usage

```
pmm2_nonlinear_onestep(
  theta_classical,
  fn_residuals,
  fn_jacobian = NULL,
  verbose = FALSE
)
```

Arguments

| | |
|-----------------|---|
| theta_classical | Parameter estimates from a classical method (e.g., MLE) |
| fn_residuals | Function(theta) returning the residual vector |
| fn_jacobian | Function(theta) returning the Jacobian matrix. If NULL, numerical Jacobian via numDeriv is used |
| verbose | Print progress |

Value

List with results (theta, residuals, etc.)

pmm2_variance_factor *Calculate theoretical skewness, kurtosis coefficients and variance reduction factor*

Description

Calculate theoretical skewness, kurtosis coefficients and variance reduction factor

Usage

```
pmm2_variance_factor(m2, m3, m4)
```

Arguments

m2, m3, m4 central moments of second, third and fourth orders

Value

List with fields c3, c4 and g

pmm2_variance_matrices *Calculate theoretical variance matrices for OLS and PMM2*

Description

Calculate theoretical variance matrices for OLS and PMM2

Usage

```
pmm2_variance_matrices(X, m2, m3, m4)
```

Arguments

`X` Design matrix with column of ones
`m2, m3, m4` central moments of OLS residuals

Value

List with fields `ols`, `pmm2`, `c3`, `c4`, `g`

| | |
|---------------|---|
| PMM3fit-class | <i>S4 class for storing PMM3 regression model results</i> |
|---------------|---|

Description

PMM3 (S=3) is designed for symmetric platykurtic error distributions. This class is fully standalone and does NOT inherit from BasePMM2.

Slots

`coefficients` numeric vector of estimated parameters
`residuals` numeric vector of final residuals
`m2` numeric second central moment of initial residuals
`m4` numeric fourth central moment of initial residuals
`m6` numeric sixth central moment of initial residuals
`gamma4` numeric excess kurtosis coefficient
`gamma6` numeric sixth-order cumulant coefficient
`g_coefficient` numeric theoretical variance reduction factor g_3
`kappa` numeric moment ratio used in NR solver
`convergence` logical indicating whether algorithm converged
`iterations` numeric number of iterations performed
`call` original function call

pmm3_variance_factor *Calculate PMM3 theoretical variance reduction factor*

Description

Computes the standardised cumulant coefficients `gamma4` and `gamma6` from the raw central moments, and derives the PMM3 efficiency factor $g_3 = 1 - \gamma_4^2 / (6 + 9\gamma_4 + \gamma_6)$.

Usage

```
pmm3_variance_factor(m2, m4, m6)
```

Arguments

| | |
|-----------------|-----------------------|
| <code>m2</code> | Second central moment |
| <code>m4</code> | Fourth central moment |
| <code>m6</code> | Sixth central moment |

Value

A list with components:

| | |
|---------------------|---------------------------------------|
| <code>gamma4</code> | Excess kurtosis |
| <code>gamma6</code> | Sixth-order cumulant coefficient |
| <code>g3</code> | Theoretical variance reduction factor |

pmm_dispatch *Automatic PMM method selection*

Description

Analyses OLS residual cumulants to recommend the best estimation method: OLS (Gaussian errors), PMM2 (asymmetric errors), or PMM3 (symmetric platykurtic errors).

Usage

```
pmm_dispatch(  
  residuals,  
  symmetry_threshold = 0.3,  
  kurtosis_threshold = -0.7,  
  g2_threshold = 0.95,  
  verbose = TRUE  
)
```

Arguments

| | |
|--------------------|--|
| residuals | numeric vector of OLS residuals |
| symmetry_threshold | numeric: gamma3 threshold for symmetry (default 0.3) |
| kurtosis_threshold | numeric: gamma4 threshold for PMM3 (default -0.7) |
| g2_threshold | numeric: minimum g2 improvement to justify PMM2 (default 0.95) |
| verbose | logical: print decision reasoning (default TRUE) |

Value

A list with components:

| | |
|-----------------|--|
| method | Character: "OLS", "PMM2", or "PMM3" |
| gamma3 | Sample skewness |
| gamma4 | Sample excess kurtosis |
| gamma6 | Sample 6th cumulant coefficient |
| g2 | PMM2 efficiency factor |
| g3 | PMM3 efficiency factor |
| g_selected | Efficiency factor for chosen method |
| improvement_pct | Expected variance reduction percentage |
| reasoning | Human-readable decision explanation |
| n | Sample size |

Examples

```
set.seed(42)
x <- rnorm(200); eps <- runif(200, -1, 1)
y <- 1 + 2 * x + eps
fit_ols <- lm(y ~ x)
pmm_dispatch(residuals(fit_ols))
```

pmm_gamma6

Compute sixth-order cumulant coefficient gamma6

Description

Calculates $\gamma_6 = m_6/m_3^3 - 15m_4/m_2^2 + 30$ from a numeric vector. For a Gaussian distribution gamma6 equals zero.

Usage

```
pmm_gamma6(x)
```

Arguments

x numeric vector

Value

Numeric scalar: the sixth-order cumulant coefficient

pmm_kurtosis *Calculate kurtosis from data*

Description

Calculate kurtosis from data

Usage

```
pmm_kurtosis(x, excess = TRUE)
```

Arguments

x numeric vector
excess logical, whether to return excess kurtosis (kurtosis - 3)

Value

Kurtosis value

pmm_skewness *Calculate skewness from data*

Description

Calculate skewness from data

Usage

```
pmm_skewness(x)
```

Arguments

x numeric vector

Value

Skewness value

 predict, PMM2fit-method

Prediction method for PMM2fit objects

Description

Computes predictions for new data using a fitted PMM2 model. The method extracts the formula from the fitted model, constructs a design matrix from the new data, and computes predictions via matrix multiplication. This approach works with arbitrary variable names and model specifications.

Usage

```
## S4 method for signature 'PMM2fit'
predict(object, newdata = NULL, debug = FALSE, ...)
```

Arguments

| | |
|---------|--|
| object | PMM2fit object returned by <code>lm_pmm2()</code> |
| newdata | Data frame containing the predictor variables with the same names as those used in the original model fit. Required parameter. |
| debug | Logical value indicating whether to output debug information about the prediction process. Default is FALSE. |
| ... | Additional arguments (currently not used) |

Details

The prediction is computed as $X\beta$ where X is the design matrix constructed from `newdata` using the model formula, and β are the estimated coefficients. The method automatically handles:

- Models with intercepts and without
- Arbitrary variable names (not limited to "x1", "x2", etc.)
- Interaction terms and transformations specified in the formula
- Automatic coefficient reordering to match design matrix columns

Value

Numeric vector of predicted values for the observations in `newdata`

Examples

```
# Fit model
fit <- lm_pmm2(mpg ~ wt + hp, data = mtcars)

# Predict on new data
newdata <- data.frame(wt = c(2.5, 3.0), hp = c(100, 150))
predictions <- predict(fit, newdata = newdata)
```

predict,PMM3fit-method

Predict method for PMM3fit objects

Description

Predict method for PMM3fit objects

Usage

```
## S4 method for signature 'PMM3fit'
predict(object, newdata = NULL, ...)
```

Arguments

| | |
|---------|-------------------------------------|
| object | PMM3fit object |
| newdata | Data frame with predictor variables |
| ... | Additional arguments (not used) |

Value

Numeric vector of predicted values

predict,TS2fit-method *Prediction method for TS2fit objects*

Description

Prediction method for TS2fit objects

Usage

```
## S4 method for signature 'TS2fit'
predict(object, n.ahead = 1, ...)
```

Arguments

| | |
|---------|--------------------------------------|
| object | TS2fit object |
| n.ahead | Number of steps ahead for prediction |
| ... | additional arguments (not used) |

Value

Vector or list of predictions, depending on model type

predict,TS3fit-method *Predict method for TS3fit objects*

Description

Predict method for TS3fit objects

Usage

```
## S4 method for signature 'TS3fit'  
predict(object, n.ahead = 1, ...)
```

Arguments

| | |
|---------|--|
| object | TS3fit object |
| n.ahead | Integer: number of steps ahead to forecast |
| ... | Additional arguments (not used) |

Value

Numeric vector of predicted values

residuals,PMM2fit-method
Extract residuals from PMM2fit object

Description

Extract residuals from PMM2fit object

Usage

```
## S4 method for signature 'PMM2fit'  
residuals(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM2fit object |
| ... | Additional arguments (not used) |

Value

Vector of residuals

residuals,PMM3fit-method

Extract residuals from PMM3fit object

Description

Extract residuals from PMM3fit object

Usage

```
## S4 method for signature 'PMM3fit'  
residuals(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM3fit object |
| ... | Additional arguments (not used) |

Value

Vector of residuals

residuals,TS2fit-method

Extract residuals from TS2fit object

Description

Extract residuals from TS2fit object

Usage

```
## S4 method for signature 'TS2fit'  
residuals(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS2fit object |
| ... | Additional arguments (not used) |

Value

Vector of residuals (innovations)

 residuals,TS3fit-method

Extract residuals from TS3fit object

Description

Extract residuals from TS3fit object

Usage

```
## S4 method for signature 'TS3fit'
residuals(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS3fit object |
| ... | Additional arguments (not used) |

Value

Vector of residuals

 SARIMAPMM2-class

S4 class for Seasonal ARIMA model results with PMM2

Description

This class stores the results of fitting a Seasonal Autoregressive Integrated Moving Average (SARIMA) model using the PMM2 method. It extends SARMA with differencing operators.

Details

The SARIMAPMM2 class represents fitted SARIMA(p,d,q) x (P,D,Q)_s models:

$$(1 - \phi_1 B - \dots - \phi_p B^p)(1 - \Phi_1 B^s - \dots - \Phi_P B^{Ps})(1 - B)^d (1 - B^s)^D y_t = (1 + \theta_1 B + \dots + \theta_q B^q)(1 + \Theta_1 B^s + \dots + \Theta_Q B^{Qs}) \epsilon_t.$$

Where B is the backshift operator.

Slots

coefficients Numeric vector of estimated parameters
 residuals Numeric vector of residuals/innovations
 m2 Second central moment of residuals
 m3 Third central moment of residuals
 m4 Fourth central moment of residuals
 convergence Logical, whether PMM2 algorithm converged
 iterations Integer, number of iterations performed
 call Original function call
 model_type Character, model type identifier ("sarima")
 intercept Numeric, intercept/mean term
 original_series Numeric vector, original time series data
 order List with model specification: list(ar, sar, ma, sma, d, D, period)

- ar: Non-seasonal AR order (p)
- sar: Seasonal AR order (P)
- ma: Non-seasonal MA order (q)
- sma: Seasonal MA order (Q)
- d: Non-seasonal differencing order
- D: Seasonal differencing order
- period: Seasonal period (s)

See Also

[sarima_pmm2](#) for fitting SARIMA models

sarima_pmm2

Fit a Seasonal ARIMA model using PMM2 method

Description

Fits a Seasonal Autoregressive Integrated Moving Average (SARIMA) model with both non-seasonal and seasonal differencing operators.

Usage

```

sarima_pmm2(
  x,
  order = c(0, 1, 0, 1),
  seasonal = list(order = c(1, 1), period = 12),
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  include.mean = NULL,

```

```

max_iter = 50,
tol = 1e-06,
regularize = TRUE,
reg_lambda = 1e-08,
ma_method = c("mle", "pmm2"),
verbose = FALSE,
multiplicative = TRUE
)

```

Arguments

| | |
|----------------|--|
| x | Numeric vector of time series data |
| order | Vector of length 4: c(p, P, q, Q) where: <ul style="list-style-type: none"> • p: Non-seasonal AR order • P: Seasonal AR order • q: Non-seasonal MA order • Q: Seasonal MA order |
| seasonal | List with seasonal specification: list(order = c(d, D), period = s) <ul style="list-style-type: none"> • d: Non-seasonal differencing order • D: Seasonal differencing order • period: Seasonal period (s) |
| method | Estimation method: "pmm2" (default), "css-ml", "css" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson, recommended for SARIMA), or "linearized" (specialized for MA/SMA models). |
| include.mean | Logical, include drift term (default TRUE for d+D=0, FALSE otherwise) |
| max_iter | Maximum iterations for PMM2 algorithm (default 50) |
| tol | Convergence tolerance (default 1e-6) |
| regularize | Logical, use regularization for numerical stability (default TRUE) |
| reg_lambda | Regularization parameter (default 1e-8) |
| ma_method | Method for MA/SMA estimation: "mle" (default) or "pmm2" |
| verbose | Logical, print progress information (default FALSE) |
| multiplicative | Logical, use multiplicative seasonal model form with cross-terms between non-seasonal and seasonal components (default TRUE). If FALSE, uses additive form. |

Details

The SARIMA(p,d,q) x (P,D,Q)_s model satisfies

$$(1 - \phi_1 B - \dots - \phi_p B^p)(1 - \Phi_1 B^s - \dots - \Phi_P B^{Ps})(1 - B)^d(1 - B^s)^D y_t = (1 + \theta_1 B + \dots + \theta_q B^q)(1 + \Theta_1 B^s + \dots + \Theta_Q B^{Qs}) \epsilon_t.$$

Where B is the backshift operator. The model combines:

- Non-seasonal differencing: $(1-B)^d$
- Seasonal differencing: $(1-B^s)^D$
- Non-seasonal ARMA components
- Seasonal ARMA components

Variant Recommendations for SARIMA:

- "unified_iterative" (recommended): Monte Carlo experiments showed 16.4% seasonal dynamics
- "unified_global" (default): Faster alternative with good accuracy
- "linearized": Not recommended for SARIMA (designed for MA/SMA)

Value

S4 object of class SARIMAPMM2 containing:

- coefficients: Estimated parameters
- residuals: Model residuals/innovations (padded with zeros to match original length)
- m2, m3, m4: Central moments of residuals
- convergence: Convergence status
- iterations: Number of iterations performed

References

Monte Carlo validation (R=50, n=200): Unified Iterative achieved 16.4% improvement for SARIMA models. See NEWS.md (Version 0.2.0).

See Also

[sarma_pmm2](#), [arima_pmm2](#)

Examples

```
set.seed(123)
n <- 200
y <- arima.sim(n = n,
  model = list(order = c(1, 0, 1), ar = 0.5, ma = 0.3,
    seasonal = list(order = c(1, 0, 0), ar = 0.4, period = 12)))
fit <- sarima_pmm2(y,
  order = c(1, 0, 1, 0),
  seasonal = list(order = c(1, 0), period = 12))
summary(fit)
```

SARMAPMM2-class

S4 class for Seasonal ARMA model results with PMM2

Description

This class stores the results of fitting a Seasonal Autoregressive Moving Average (SARMA) model using the PMM2 method. It combines both seasonal AR and seasonal MA components.

Details

The SARMAPMM2 class represents fitted SARMA models combining:

- AR(p): $\phi_1 y_{t-1} + \dots + \phi_p y_{t-p}$
- Seasonal AR component: $\Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps}$
- MA(q): $\theta_1 \epsilon_{t-1} + \dots + \theta_q \epsilon_{t-q}$
- Seasonal MA component: $\Theta_1 \epsilon_{t-s} + \dots + \Theta_Q \epsilon_{t-Qs}$

Slots

`coefficients` Numeric vector of estimated parameters (SAR and SMA coefficients)

`residuals` Numeric vector of residuals/innovations

`m2` Second central moment of residuals

`m3` Third central moment of residuals

`m4` Fourth central moment of residuals

`convergence` Logical, whether PMM2 algorithm converged

`iterations` Integer, number of iterations performed

`call` Original function call

`model_type` Character, model type identifier ("sarma")

`intercept` Numeric, intercept/mean term

`original_series` Numeric vector, original time series data

`order` List with model specification: `list(ar, sar, ma, sma, period)`

- `ar`: Non-seasonal AR order (p)
- `sar`: Seasonal AR order (P)
- `ma`: Non-seasonal MA order (q)
- `sma`: Seasonal MA order (Q)
- `period`: Seasonal period (s)

See Also

[sarma_pmm2](#) for fitting SARMA models

 sarma_pmm2

Fit a Seasonal ARMA model using PMM2 method

Description

Fits a Seasonal Autoregressive Moving Average (SARMA) model that combines both seasonal AR and seasonal MA components, optionally with non-seasonal AR and MA terms as well.

Usage

```
sarma_pmm2(
  x,
  order = c(0, 1, 0, 1),
  season = list(period = 12),
  method = "pmm2",
  include.mean = TRUE,
  max_iter = 50,
  tol = 1e-06,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Vector of length 4: c(p, P, q, Q) where: <ul style="list-style-type: none"> • p: Non-seasonal AR order • P: Seasonal AR order • q: Non-seasonal MA order • Q: Seasonal MA order |
| season | List with seasonal specification: list(period = s) where s is the seasonal period (e.g., 12 for monthly data) |
| method | Estimation method: "pmm2" (default), "css-ml", "css" |
| include.mean | Logical, include intercept/mean term (default TRUE) |
| max_iter | Maximum iterations for PMM2 algorithm (default 50) |
| tol | Convergence tolerance (default 1e-6) |
| regularize | Logical, use regularization for numerical stability (default TRUE) |
| reg_lambda | Regularization parameter (default 1e-8) |
| verbose | Logical, print progress information (default FALSE) |

Details

The SARMA model combines four components:

- AR(p): $\phi_1 y_{t-1} + \dots + \phi_p y_{t-p}$
- Seasonal AR: $\Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps}$
- MA(q): $\theta_1 \epsilon_{t-1} + \dots + \theta_q \epsilon_{t-q}$
- Seasonal MA: $\Theta_1 \epsilon_{t-s} + \dots + \Theta_Q \epsilon_{t-Qs}$

The PMM2 method provides more efficient parameter estimates than ML/CSS when the innovation distribution is asymmetric (non-Gaussian).

Value

S4 object of class SARMAPMM2 containing:

- coefficients: Estimated parameters
- residuals: Model residuals/innovations (padded with zeros to match original length)
- m2, m3, m4: Central moments of residuals
- convergence: Convergence status
- iterations: Number of iterations performed

See Also

[sar_pmm2](#), [sma_pmm2](#), [sarima_pmm2](#)

Examples

```
# Generate synthetic seasonal data with SARMA structure
set.seed(123)
n <- 200
y <- arima.sim(n = n, list(
  ar = 0.5, ma = 0.3,
  seasonal = list(sar = 0.6, sma = 0.4, period = 12)
))

# Fit SARMA(1,1,1,1)_12 model with PMM2
fit <- sarma_pmm2(y, order = c(1, 1, 1, 1), season = list(period = 12))
summary(fit)

# Pure seasonal model (no non-seasonal components)
fit_pure <- sarma_pmm2(y, order = c(0, 1, 0, 1), season = list(period = 12))
```

SARPMM2-class

*S4 class for Seasonal AR model results with PMM2***Description**

This class stores the results of fitting a Seasonal Autoregressive (SAR) model using the PMM2 method. It extends the TS2fit class with additional slots specific to seasonal models.

Details

The SARPMM2 class represents fitted SAR models of the form

$$y_t = \phi_1 y_{t-1} + \cdots + \phi_p y_{t-p} + \Phi_1 y_{t-s} + \cdots + \Phi_P y_{t-Ps} + \epsilon_t.$$

Where:

- p is the non-seasonal AR order
- P is the seasonal AR order
- s is the seasonal period

Slots

`coefficients` Numeric vector of estimated parameters (AR and SAR coefficients)

`residuals` Numeric vector of residuals/innovations

`m2` Second central moment of residuals

`m3` Third central moment of residuals

`m4` Fourth central moment of residuals

`convergence` Logical, whether PMM2 algorithm converged

`iterations` Integer, number of iterations performed

`call` Original function call

`model_type` Character, model type identifier ("sar")

`intercept` Numeric, intercept/mean term

`original_series` Numeric vector, original time series data

`order` List with model specification: list(ar, sar, period)

- ar: Non-seasonal AR order (p)
- sar: Seasonal AR order (P)
- period: Seasonal period (s)

See Also

[sar_pmm2](#) for fitting SAR models

sar_pmm2

*Fit Seasonal AR model using PMM2 method***Description**

Fits a Seasonal Autoregressive (SAR) model using the Polynomial Maximization Method (PMM2). The model can include both non-seasonal and seasonal AR components.

Usage

```

sar_pmm2(
  x,
  order = c(0, 1),
  season = list(period = 12),
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  include.mean = TRUE,
  multiplicative = FALSE,
  max_iter = 50,
  tol = 1e-06,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)

```

Arguments

| | |
|----------------|--|
| x | Numeric vector of time series data |
| order | Vector of length 2: c(p, P) where: <ul style="list-style-type: none"> • p: Non-seasonal AR order • P: Seasonal AR order |
| season | List with seasonal specification: list(period = s) where s is the seasonal period (e.g., 12 for monthly data with annual seasonality) |
| method | Estimation method: "pmm2" (default), "ols", "css" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction from MLE/CSS), "unified_iterative" (full Newton-Raphson), or "linearized" (not recommended for SAR models). |
| include.mean | Logical, include intercept/mean term (default TRUE) |
| multiplicative | Logical, use multiplicative form with cross-terms (default FALSE) |
| max_iter | Maximum iterations for PMM2 algorithm (default 50) |
| tol | Convergence tolerance (default 1e-6) |
| regularize | Logical, use regularization for numerical stability (default TRUE) |
| reg_lambda | Regularization parameter (default 1e-8) |
| verbose | Logical, print progress information (default FALSE) |

Details

The SAR model has the form

$$y_t = \phi_1 y_{t-1} + \dots + \phi_p y_{t-p} + \Phi_1 y_{t-s} + \dots + \Phi_P y_{t-Ps} + \mu + \epsilon_t.$$

Where:

- p is the non-seasonal AR order
- P is the seasonal AR order
- s is the seasonal period
- ϵ_t are innovations (errors)

The PMM2 method provides more efficient parameter estimates than OLS when the innovation distribution is asymmetric (non-Gaussian). The expected variance reduction is given by $g = 1 - c_3^2 / (2 + c_4)$, where c_3 and c_4 are the skewness and excess kurtosis coefficients.

Variant Recommendations for SAR:

- "unified_global" (default): Fast one-step correction, suitable for most SAR models
- "unified_iterative": Best accuracy for complex seasonal patterns
- "linearized": Not recommended for SAR (designed for MA/SMA)

Value

S4 object of class SARPMM2 containing:

- coefficients: Estimated AR and SAR parameters
- residuals: Model residuals/innovations (padded with zeros to match original length)
- m2, m3, m4: Central moments of residuals
- convergence: Convergence status
- iterations: Number of iterations performed

See Also

[ar_pmm2](#), [ts_pmm2](#), [compare_sar_methods](#)

Examples

```
# Generate synthetic seasonal data
n <- 120
y <- arima.sim(n = n, list(ar = 0.7, seasonal = list(sar = 0.5, period = 12)))

# Fit SAR(1,1)_12 model with PMM2
fit <- sar_pmm2(y, order = c(1, 1), season = list(period = 12))
summary(fit)

# Simple seasonal model (no non-seasonal component)
fit_pure_sar <- sar_pmm2(y, order = c(0, 1), season = list(period = 12))
```

```
# Compare with OLS
fit_ols <- sar_pmm2(y, order = c(1, 1), season = list(period = 12), method = "ols")
```

SMAPMM2-class

S4 class for Seasonal MA PMM2 results

Description

This class stores the results of fitting a Seasonal Moving Average (SMA) model using the Polynomial Maximization Method (PMM2).

Details

The SMA(Q)_s model is expressed as

$$y_t = \mu + \epsilon_t + \Theta_1 \epsilon_{t-s} + \Theta_2 \epsilon_{t-2s} + \dots + \Theta_Q \epsilon_{t-Qs}.$$

Where:

- Q is the seasonal MA order
- s is the seasonal period
- epsilon_t are innovations

Slots

coefficients Estimated seasonal MA coefficients (Theta_1, Theta_2, ..., Theta_Q)

innovations Estimated innovations (residuals epsilon_t)

m2 Second central moment (variance) of innovations

m3 Third central moment (skewness indicator) of innovations

m4 Fourth central moment (kurtosis indicator) of innovations

convergence Logical indicating whether PMM2 algorithm converged

iterations Number of iterations required for convergence

call The function call that created this object

model_type Character string "sma"

intercept Model intercept (mean)

original_series Original time series data

order List with Q (seasonal MA order) and s (seasonal period)

See Also

[sma_pmm2](#) for fitting SMA models

sma_pmm2

*Fit a Seasonal MA model using PMM2***Description**

Fits a Seasonal Moving Average (SMA) model using the Polynomial Maximization Method (PMM2). This is particularly effective when the innovations have asymmetric or non-Gaussian distributions.

Usage

```
sma_pmm2(
  x,
  order = 1,
  season = list(period = 12),
  method = "pmm2",
  pmm2_variant = c("unified_global", "unified_iterative", "linearized"),
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  na.action = na.fail,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector (time series data) |
| order | Seasonal MA order (Q) |
| season | List with seasonal specification: list(period = s) |
| method | Estimation method: "pmm2" or "css" |
| pmm2_variant | Character string specifying PMM2 implementation variant. Options: "unified_global" (default, one-step correction), "unified_iterative" (full Newton-Raphson), or "linearized" (specialized for MA/SMA models, recommended for SMA). |
| max_iter | Maximum iterations for PMM2 algorithm |
| tol | Convergence tolerance |
| include.mean | Include intercept in the model |
| na.action | Function to handle missing values |
| regularize | Add regularization to Jacobian matrix |
| reg_lambda | Regularization parameter |
| verbose | Print diagnostic information |

Details

The SMA(Q)_s model has the form

$$y_t = \mu + \epsilon_t + \Theta_1 \epsilon_{t-s} + \Theta_2 \epsilon_{t-2s} + \dots + \Theta_Q \epsilon_{t-Qs}.$$

Where:

- Q is the seasonal MA order
- s is the seasonal period (12 for monthly, 4 for quarterly)
- epsilon_t are innovations (errors)

The PMM2 method provides more efficient parameter estimates than ML/CSS when the innovation distribution is asymmetric (non-Gaussian). The expected variance reduction is given by $g = 1 - c3^2 / (2 + c4)$, where c3 and c4 are the skewness and excess kurtosis coefficients.

Variant Recommendations for SMA:

- "linearized" (recommended): Fastest and most stable for SMA models, avoids complex Jacobian computation while maintaining accuracy
- "unified_global" (default): Good balance of speed and accuracy
- "unified_iterative": Best accuracy but slower, use for complex SMA patterns

Value

An S4 object of class SMAPMM2 containing:

- coefficients: Seasonal MA coefficients (Theta_1, Theta_2, ..., Theta_Q)
- innovations: Estimated innovations (residuals)
- m2, m3, m4: Central moments of innovations
- convergence: Convergence status
- iterations: Number of iterations performed
- intercept: Model intercept
- original_series: Original time series
- order: Model order list(Q, s)

See Also

[ma_pmm2](#), [sar_pmm2](#), [arima_pmm2](#)

Examples

```
# Generate synthetic seasonal data
set.seed(123)
n <- 120
s <- 12
theta <- 0.6

# Gamma innovations (asymmetric)
```

```

innov <- rgamma(n, shape = 2, scale = 1) - 2
y <- numeric(n)
for (t in 1:n) {
  ma_term <- if (t > s) theta * innov[t - s] else 0
  y[t] <- innov[t] + ma_term
}

# Fit SMA(1)_12 model with PMM2
fit <- sma_pmm2(y, order = 1, season = list(period = 12))
summary(fit)

# Compare with CSS
fit_css <- sma_pmm2(y, order = 1, season = list(period = 12), method = "css")

```

| | |
|-----------------|-------------------------|
| solve_pmm2_step | <i>PMM2 step solver</i> |
|-----------------|-------------------------|

Description

Solves the linearized system to find the parameter update. Based on the Taylor expansion: $e(\theta) \sim e(\theta_k) - J * \delta$

Usage

```
solve_pmm2_step(residuals, J, pmm_stats)
```

Arguments

| | |
|-----------|--|
| residuals | Current residuals |
| J | Jacobian matrix (n x p), where $J_{ij} = d(\hat{y}_i)/d(\theta_j)$. We assume the standard regression definition: $y = f(\theta) + e$, so $e = y - f(\theta)$ and $d(e)/d(\theta) = -d(f)/d(\theta)$. We expect $J = d(f)/d(\theta)$ (gradient of the regression function). |
| pmm_stats | Statistics from compute_pmm2_components |

Value

Update vector delta

 summary,PMM2fit-method

Generic summary method for PMM2fit objects

Description

Generic summary method for PMM2fit objects

Usage

```
## S4 method for signature 'PMM2fit'
summary(object, formula = NULL, data = NULL, B = 100, ...)
```

Arguments

| | |
|---------|--|
| object | object of class "PMM2fit" |
| formula | (optional) formula used for the model |
| data | (optional) data used |
| B | number of bootstrap replications for statistical inference |
| ... | additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly).

summary,PMM3fit-method

Summary method for PMM3fit objects

Description

Summary method for PMM3fit objects

Usage

```
## S4 method for signature 'PMM3fit'
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | PMM3fit object |
| ... | Additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly)

summary, SARIMAPMM2-method

Generic summary method for SARIMAPMM2 objects

Description

Generic summary method for SARIMAPMM2 objects

Usage

```
## S4 method for signature 'SARIMAPMM2'
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | object of class "SARIMAPMM2" |
| ... | additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly).

summary, SARMAPMM2-method

Generic summary method for SARMAPMM2 objects

Description

Generic summary method for SARMAPMM2 objects

Usage

```
## S4 method for signature 'SARMAPMM2'
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | object of class "SARMAPMM2" |
| ... | additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly).

summary, SARPMM2-method

Summary method for SARPMM2 objects

Description

Summary method for SARPMM2 objects

Usage

```
## S4 method for signature 'SARPMM2'  
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | SARPMM2 object |
| ... | Additional arguments (not used) |

Value

Invisibly returns the object

summary, SMAPMM2-method

Summary method for SMAPMM2 objects

Description

Summary method for SMAPMM2 objects

Usage

```
## S4 method for signature 'SMAPMM2'  
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | SMAPMM2 object |
| ... | Additional arguments (not used) |

Value

Invisibly returns the object

summary,TS2fit-method *Generic summary method for TS2fit objects*

Description

Generic summary method for TS2fit objects

Usage

```
## S4 method for signature 'TS2fit'  
summary(object, ...)
```

Arguments

| | |
|--------|--------------------------------------|
| object | object of class "TS2fit" or subclass |
| ... | additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly).

summary,TS3fit-method *Summary method for TS3fit objects*

Description

Summary method for TS3fit objects

Usage

```
## S4 method for signature 'TS3fit'  
summary(object, ...)
```

Arguments

| | |
|--------|---------------------------------|
| object | TS3fit object |
| ... | Additional arguments (not used) |

Value

Prints summary to console; returns object (invisibly)

| | |
|---------------|---|
| test_symmetry | <i>Test whether residuals are sufficiently symmetric for PMM3</i> |
|---------------|---|

Description

Computes the skewness coefficient γ_3 and checks whether its absolute value falls below a given threshold. This helps decide between PMM2 (asymmetric) and PMM3 (symmetric platykurtic) estimation.

Usage

```
test_symmetry(x, threshold = 0.3)
```

Arguments

| | |
|-----------|--|
| x | numeric vector of residuals |
| threshold | numeric threshold for $ \gamma_3 $ (default 0.3) |

Value

A list with components:

| | |
|--------------|---|
| gamma3 | Sample skewness coefficient |
| is_symmetric | Logical: TRUE if $ \gamma_3 \leq \text{threshold}$ |
| message | Human-readable verdict |

| | |
|--------------|---|
| TS2fit-class | <i>Base S4 class for storing PMM2 time series model results</i> |
|--------------|---|

Description

Base class for storing results of time series model estimation using PMM2

Slots

| | |
|--------------|--|
| coefficients | numeric vector of estimated parameters |
| residuals | numeric vector of final residuals |
| m2 | numeric second central moment of initial residuals |
| m3 | numeric third central moment of initial residuals |
| m4 | numeric fourth central moment of initial residuals |
| convergence | logical or integer code indicating whether algorithm converged |
| iterations | numeric number of iterations performed |
| call | original function call |

model_type character string indicating model type
intercept numeric value of intercept
original_series numeric vector of original time series
order list of order parameters

Slots

coefficients Estimated coefficients
residuals Final residuals
m2 Second central moment
m3 Third central moment
m4 Fourth central moment
convergence Convergence status
iterations Number of iterations performed
call Original call
model_type Model type
intercept Intercept
original_series Original time series
order Model orders

TS3fit-class

Base S4 class for PMM3 time series model results

Description

Stores results from time series estimation using PMM3 (S=3). Designed for symmetric platykurtic innovations. Does NOT inherit from BasePMM2 or TS2fit.

Slots

coefficients numeric vector of estimated parameters
residuals numeric vector of final residuals/innovations
m2 numeric second central moment of initial residuals
m4 numeric fourth central moment of initial residuals
m6 numeric sixth central moment of initial residuals
gamma4 numeric excess kurtosis coefficient
gamma6 numeric sixth-order cumulant coefficient
g_coefficient numeric theoretical variance reduction factor g3
kappa numeric moment ratio used in NR solver
convergence logical whether algorithm converged

iterations numeric number of iterations performed
 call original function call
 model_type character string indicating model type
 intercept numeric intercept value
 original_series numeric vector of original time series
 order list of order parameters

 ts_pmm2

Fit a time series model using the PMM2 method

Description

Fit a time series model using the PMM2 method

Usage

```

ts_pmm2(
  x,
  order,
  model_type = c("ar", "ma", "arma", "arima"),
  method = "pmm2",
  max_iter = 50,
  tol = 1e-06,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  regularize = TRUE,
  reg_lambda = 1e-08,
  verbose = FALSE
)

```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification: - For AR models: a single integer (AR order) - For MA models: a single integer (MA order) - For ARMA models: vector c(p, q) (AR and MA orders) - For ARIMA models: vector c(p, d, q) (AR, differencing, and MA orders) |
| model_type | String specifying the model type: "ar", "ma", "arma", or "arima" |
| method | String: estimation method, one of "pmm2" (default), "css", "ml", "yw", "ols" |
| max_iter | Integer: maximum number of iterations for the algorithm |
| tol | Numeric: tolerance for convergence |
| include.mean | Logical: whether to include a mean (intercept) term |

| | |
|------------|---|
| initial | List or vector of initial parameter estimates (optional) |
| na.action | Function for handling missing values, default is na.fail |
| regularize | Logical, add small values to diagonal for numerical stability |
| reg_lambda | Regularization parameter (if regularize=TRUE) |
| verbose | Logical: whether to print progress information |

Details

The PMM2 algorithm works as follows:

1. Fits an initial model using a standard method (OLS, Yule-Walker, CSS or ML)
2. Computes central moments (m2, m3, m4) from initial residuals/innovations
3. Uses these moments with a specialized solver (pmm2_algorithm) to find robust parameter estimates

Value

An S4 object TS2fit of the corresponding subclass

| | |
|-------------------|--|
| ts_pmm2_inference | <i>Bootstrap inference for PMM2 time series models</i> |
|-------------------|--|

Description

Bootstrap inference for PMM2 time series models

Usage

```
ts_pmm2_inference(
  object,
  x = NULL,
  B = 200,
  seed = NULL,
  block_length = NULL,
  method = c("residual", "block"),
  parallel = FALSE,
  cores = NULL,
  debug = FALSE
)
```

Arguments

| | |
|--------------|---|
| object | object of class TS2fit |
| x | (optional) original time series; if NULL, uses object@original_series |
| B | number of bootstrap replications |
| seed | (optional) for reproducibility |
| block_length | block length for block bootstrap; if NULL, uses heuristic value |
| method | bootstrap type: "residual" or "block" |
| parallel | logical, whether to use parallel computing |
| cores | number of cores for parallel computing |
| debug | logical, whether to output additional diagnostic information |

Value

data.frame with columns: Estimate, Std.Error, t.value, p.value

 ts_pmm3

Fit a time series model using PMM3

Description

Core function that fits AR, MA, ARMA, or ARIMA models using the Polynomial Maximization Method of order 3 (PMM3). Designed for symmetric platykurtic innovations.

Usage

```
ts_pmm3(
  x,
  order,
  model_type = c("ar", "ma", "arma", "arima"),
  max_iter = 100,
  tol = 1e-06,
  adaptive = FALSE,
  step_max = 5,
  include.mean = TRUE,
  initial = NULL,
  na.action = na.fail,
  verbose = FALSE
)
```

Arguments

| | |
|--------------|---|
| x | Numeric vector of time series data |
| order | Model order specification (see <code>ts_pmm2</code> for format) |
| model_type | Character: "ar", "ma", "arma", or "arima" |
| max_iter | Integer: maximum NR iterations (default 100) |
| tol | Numeric: convergence tolerance (default 1e-6) |
| adaptive | Logical: re-estimate kappa each iteration (default FALSE) |
| step_max | Numeric: maximum NR step size (default 5.0) |
| include.mean | Logical: include mean/intercept term (default TRUE) |
| initial | Optional initial parameter estimates |
| na.action | Function for handling missing values (default na.fail) |
| verbose | Logical: print progress information (default FALSE) |

Details

The PMM3 time series algorithm:

1. Obtains initial estimates via classical methods (OLS/YW for AR, CSS for MA/ARMA/ARIMA)
2. Computes moments m_2 , m_4 , m_6 and kappa from initial residuals
3. Checks symmetry: warns if $|\text{lgamma}_3| > 0.3$
4. Applies Newton-Raphson with PMM3 score equations

PMM3 is beneficial when innovations are symmetric and platykurtic ($\text{gamma}_4 < 0$), e.g. uniform, truncated normal.

Value

An S4 object of the appropriate TS3fit subclass

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