

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)>, and Zabolotnii (2025) <[doi:10.48550/arXiv.2511.07059](https://doi.org/10.48550/arXiv.2511.07059)>.

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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

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

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 'pmm2_package.R' 'pmm2_ts_design.R' 'pmm2_ts_main.R'
 'pmm2_ts_methods.R' 'pmm2_ts_recursive.R' 'pmm2_unified.R'
 'pmm2_utils.R' 'pmm3_classes.R' 'pmm3_dispatch.R' 'pmm3_main.R'
 'pmm3_solver.R' 'pmm3_ts_classes.R' 'pmm3_ts_main.R'
 'pmm3_ts_methods.R' 'pmm3_utils.R' 'pmm_show_methods.R'
 'pmm_unified_api.R' 'sarimax_wrapper.R'

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

S4 class for storing PMM2 ARIMA model results

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

S4 class for PMM3 ARIMA model results

arima_pmm2	<i>Fit an ARIMA model using PMM2 (wrapper)</i>
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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"),  
  ma_solver = c("linearized", "recursive"),  
  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).
ma_solver	String: how the moving-average recursion is treated, "linearized" (default) or "recursive". Has no effect when the MA order is zero. See Details.
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

x	Numeric vector of time series data
order	Numeric vector c(p, d, q): AR, differencing, and MA orders
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 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>
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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

*Fit an ARMA model using PMM2 (wrapper)***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"),
  ma_solver = c("linearized", "recursive"),
  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).
ma_solver	String: how the moving-average recursion is treated, "linearized" (default) or "recursive". Has no effect when the MA order is zero. See Details.
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

<code>initial</code>	List or vector of initial parameter estimates (optional)
<code>na.action</code>	Function for handling missing values, default is <code>na.fail</code>
<code>regularize</code>	Logical, add small values to diagonal for numerical stability
<code>reg_lambda</code>	Regularization parameter (if <code>regularize=TRUE</code>)
<code>verbose</code>	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

x	Numeric vector of time series data
order	Numeric vector c(p, q): AR and MA orders
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 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	<i>S4 class for storing PMM2 AR model results</i>
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Description

S4 class for storing PMM2 AR model results

ARPM3-class	<i>S4 class for PMM3 AR model results</i>
-------------	---

Description

S4 class for PMM3 AR model results

ar_pmm2	<i>Fit an AR model using PMM2 (wrapper)</i>
---------	---

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

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 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 PMM2 on asymmetric regression residuals and to illustrate dispatcher diagnostics on practical automotive data.

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

Practical regression examples:

- **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** (diagnostic, quadratic): residuals are nearly symmetric but have positive excess kurtosis, so the current `pmm_dispatch()` rule does not treat this as a PMM3 platykurtic case.

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](https://doi.org/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

Virtual S4 class for the PMM2 model family

Description

Virtual intermediate that inherits common slots (coefficients, residuals, convergence, iterations, call) from [PMMfit](#) and adds the PMM2-specific central moments m_2, m_3, m_4 . Concrete subclasses are [PMM2fit](#) (regression) and the time-series classes under [TS2fit](#).

Slots

m2 numeric second central moment of initial residuals
m3 numeric third central moment of initial residuals
m4 numeric fourth central moment of initial residuals

BasePMM3-class

Virtual S4 class for the PMM3 model family

Description

Virtual intermediate that inherits common slots from [PMMfit](#) and adds the PMM3-specific moments and cumulant coefficients used for symmetric platykurtic errors. Concrete subclasses are [PMM3fit](#) (regression) and the time-series classes under [TS3fit](#).

Slots

- 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 by the Newton-Raphson solver

coef,PMM2fit-method	<i>Extract coefficients from PMM2fit object</i>
---------------------	---

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	<i>Extract coefficients from PMM3fit object</i>
---------------------	---

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	<i>Extract coefficients from SARPMM2 object</i>
----------------------	---

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	<i>Extract coefficients from SMAPMM2 object</i>
----------------------	---

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	<i>Extract coefficients from TS2fit object</i>
--------------------	--

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	<i>Extract coefficients from TS3fit object</i>
--------------------	--

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_arima_methods` *Compare ARIMA methods*

Description

Compare ARIMA methods

Usage

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

Arguments

<code>x</code>	Numeric vector of time series data
<code>order</code>	Model order specification (see <code>ts_pmm2</code> for format)
<code>include.mean</code>	Logical, whether to include intercept term
<code>pmm2_args</code>	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_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

<code>x</code>	Numeric vector of time series data
<code>order</code>	Model order specification (see <code>ts_pmm2</code> for format)
<code>include.mean</code>	Logical, whether to include intercept term
<code>pmm2_args</code>	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).

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

Description

Compare AR methods

Usage

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

Arguments

<code>x</code>	Numeric vector of time series data
<code>order</code>	Model order specification (see <code>ts_pmm2</code> for format)
<code>include.mean</code>	Logical, whether to include intercept term
<code>pmm2_args</code>	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_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", "arma"),
  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 "arma"
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	<i>Compute central moments for PMM3 from residuals</i>
----------------------	--

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)

confint,PMM2fit-method

Confidence intervals for PMM2fit coefficients

Description

Computes normal-approximation confidence intervals using the asymptotic covariance matrix from [vcov,PMM2fit-method](#).

Usage

```
## S4 method for signature 'PMM2fit'
confint(object, parm, level = 0.95, ...)
```

Arguments

object	PMM2fit object
parm	character or integer vector of parameter names/indices to include
level	confidence level (default 0.95)
...	Additional arguments (not used)

Value

Matrix with lower and upper confidence limits

 confint, PMM3fit-method

Confidence intervals for PMM3fit coefficients

Description

Computes normal-approximation confidence intervals using the asymptotic covariance matrix from [vcov, PMM3fit-method](#).

Usage

```
## S4 method for signature 'PMM3fit'
confint(object, parm, level = 0.95, ...)
```

Arguments

object	PMM3fit object
parm	character or integer vector of parameter names/indices to include
level	confidence level (default 0.95)
...	Additional arguments (not used)

Value

Matrix with lower and upper confidence limits

 confint, TS2fit-method *Confidence intervals for TS2fit AR model coefficients*

Description

For AR models, computes normal-approximation CIs using [vcov, TS2fit-method](#). For MA/ARMA/ARIMA models, use [ts_pmm2_inference](#) instead.

Usage

```
## S4 method for signature 'TS2fit'
confint(object, parm, level = 0.95, ...)
```

Arguments

object	TS2fit (or subclass) object
parm	character or integer vector of parameter names/indices
level	confidence level (default 0.95)
...	Additional arguments (not used)

Value

Matrix with lower and upper confidence limits

confint, TS3fit-method *Confidence intervals for TS3fit AR model coefficients*

Description

For AR models, computes normal-approximation CIs using `vcov, TS3fit-method`. For MA/ARMA/ARIMA models, refit the analogous PMM2 model and use `ts_pmm2_inference`.

Usage

```
## S4 method for signature 'TS3fit'
confint(object, parm, level = 0.95, ...)
```

Arguments

object	TS3fit (or subclass) object
parm	character or integer vector of parameter names/indices
level	confidence level (default 0.95)
...	Additional arguments (not used)

Value

Matrix with lower and upper confidence limits

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	<i>Create design matrix for seasonal AR model</i>
-------------------	---

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:

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

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

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

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

format.PMMdispatch	<i>Format method for PMMdispatch objects</i>
--------------------	--

Description

Format method for PMMdispatch objects

Usage

```
## S3 method for class 'PMMdispatch'  
format(x, ...)
```

Arguments

- x object of class PMMdispatch
- ... additional arguments (ignored)

Value

A character string summarising the selected method.

get_sarimax_jacobian	<i>Calculate SARIMAX Jacobian (Numerical)</i>
----------------------	---

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 *PMM2: Main function for PMM2 (S=2)*

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)
```

logLik.PMM2fit

Extract log-likelihood from PMM2fit object

Description

Returns a Gaussian approximate log-likelihood. Defined as an S3 method so that stats::AIC and stats::BIC (which dispatch logLik via UseMethod) work without explicit S4 methods.

Usage

```
## S3 method for class 'PMM2fit'
logLik(object, ...)
```

Arguments

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

Value

Object of class logLik

logLik.PMM3fit	<i>Extract log-likelihood from PMM3fit object</i>
----------------	---

Description

Returns a Gaussian approximate log-likelihood. Defined as an S3 method so that `stats::AIC` and `stats::BIC` (which dispatch `logLik` via `UseMethod`) work without explicit S4 methods.

Usage

```
## S3 method for class 'PMM3fit'
logLik(object, ...)
```

Arguments

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

Value

Object of class `logLik`

logLik.TS2fit	<i>Extract log-likelihood from TS2fit object</i>
---------------	--

Description

Returns a Gaussian approximate log-likelihood. Non-finite residuals (which can arise from differencing or burn-in) are dropped so that AIC/BIC dispatched through this method match the convention of `compare_ts_methods()`. Defined as an S3 method so that `stats::AIC` and `stats::BIC` dispatch correctly via `UseMethod("logLik")`.

Usage

```
## S3 method for class 'TS2fit'
logLik(object, ...)
```

Arguments

object	TS2fit (or subclass) object
...	Additional arguments (not used)

Value

Object of class `logLik`

<code>logLik.TS3fit</code>	<i>Extract log-likelihood from TS3fit object</i>
----------------------------	--

Description

Returns a Gaussian approximate log-likelihood. Defined as an S3 method so that `stats::AIC` and `stats::BIC` dispatch correctly via `UseMethod("logLik")`.

Usage

```
## S3 method for class 'TS3fit'
logLik(object, ...)
```

Arguments

<code>object</code>	TS3fit (or subclass) object
<code>...</code>	Additional arguments (not used)

Value

Object of class `logLik`

<code>MAPMM2-class</code>	<i>S4 class for storing PMM2 MA model results</i>
---------------------------	---

Description

S4 class for storing PMM2 MA model results

<code>MAPMM3-class</code>	<i>S4 class for PMM3 MA model results</i>
---------------------------	---

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"),
  ma_solver = c("linearized", "recursive"),
  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).
ma_solver	String: how the moving-average recursion is treated, "linearized" (default) or "recursive". Has no effect when the MA order is zero. See Details.
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)

<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 <code>FALSE</code>)

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)
```

<code>nobs,PMM2fit-method</code>	<i>Number of observations in PMM2fit object</i>
----------------------------------	---

Description

Number of observations in `PMM2fit` object

Usage

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

Arguments

<code>object</code>	PMM2fit object
<code>...</code>	Additional arguments (not used)

Value

Integer number of observations

nobs,PMM3fit-method	<i>Number of observations in PMM3fit object</i>
---------------------	---

Description

Number of observations in PMM3fit object

Usage

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

Arguments

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

Value

Integer number of observations

nobs,TS2fit-method	<i>Number of observations in TS2fit object</i>
--------------------	--

Description

Returns the effective sample size (length of the residual vector).

Usage

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

Arguments

object	TS2fit (or subclass) object
...	Additional arguments (not used)

Value

Integer effective sample size

nobs,TS3fit-method	<i>Number of observations in TS3fit object</i>
--------------------	--

Description

Number of observations in TS3fit object

Usage

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

Arguments

object	TS3fit (or subclass) object
...	Additional arguments (not used)

Value

Integer effective sample size

plot,PMM2fit,missing-method	<i>Plot diagnostic plots for PMM2fit object</i>
-----------------------------	---

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	<i>S4 class for storing PMM2 regression model results</i>
---------------	---

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	<i>Bootstrap inference for PMM2 fit</i>
----------------	---

Description

Residual-bootstrap standard errors, p-values, and confidence intervals for a PMM2 regression fit. Three CI methods are available (see `ci_method`).

Usage

```
pmm2_inference(
  object,
  formula,
  data,
  B = 200,
  seed = NULL,
  parallel = FALSE,
  cores = NULL,
  ci_method = c("normal", "percentile", "basic")
)
```

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
ci_method	character: confidence-interval method.

"normal" (default) Symmetric Wald interval $\hat{\beta} \pm z_{1-\alpha/2} \widehat{SE}$ using the bootstrap standard deviation as $\widehat{SE}$. Always contains the point estimate and is robust to mild finite-sample bias of the bootstrap distribution.

"percentile" Empirical 2.5/97.5 percentiles of the bootstrap re-estimates (Efron's original percentile method). Not centred on the point estimate when the bootstrap distribution is biased.

"basic" Davison & Hinkley (1997) basic/pivotal interval $[2\hat{\beta} - q_{1-\alpha/2}, 2\hat{\beta} - q_{\alpha/2}]$, reflecting the percentile bracket about the point estimate.

The returned object includes a bias column ($\text{mean}(\text{boot}) - \hat{\beta}$) so users can detect finite-sample bias and switch CI methods if desired.

Value

data.frame with columns: Estimate, Std.Error, bias, t.value, p.value, conf.low, conf.high.

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). $J_{ij} = d(y_hat_i)/d(theta_j) = -d(epsilon_i)/d(theta_j)$. 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

S4 class for PMM3 regression fit results

Description

Concrete class returned by `lm_pmm3`. Inherits all slots from `BasePMM3`, which in turn inherits common slots from `PMMfit`.

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

m2	Second central moment
m4	Fourth central moment
m6	Sixth central moment

Value

A list with components:

gamma4	Excess kurtosis
gamma6	Sixth-order cumulant coefficient
g3	Theoretical variance reduction factor

pmm3_variance_matrices *Calculate theoretical variance matrices for OLS and PMM3*

Description

Returns the asymptotic covariance matrices $V_{OLS} = m_2(X^\top X)^{-1}$ and $V_{PMM3} = g_3 m_2(X^\top X)^{-1}$ where $g_3 = 1 - \gamma_4^2 / (6 + 9\gamma_4 + \gamma_6)$.

Usage

```
pmm3_variance_matrices(X, m2, m4, m6)
```

Arguments

X	Design matrix with column of ones for the intercept.
m2, m4, m6	Second, fourth, and sixth central moments computed from the initial OLS residuals.

Details

The PMM3 formula is the symmetric-platykurtic specialisation of Kunchenko's polynomial-maximisation framework ($s = 3$); see `notes/pmm3_vcov_derivation.md` for the derivation and Zabolotnii et al. (2018, 2022, 2023) for the general method. Under Gaussian errors $g_3 = 1$ and the matrix coincides with the OLS covariance.

Value

A list with components `ols` (OLS covariance matrix), `pmm3` (PMM3 covariance matrix), and the scalar efficiency diagnostics `gamma4`, `gamma6`, `g3`.

PMMfit-class

Virtual root S4 class for PMM fit objects

Description

Common parent of every concrete fit class returned by the package (PMM2fit, PMM3fit and the time-series subclasses). Holds the slots that are meaningful regardless of polynomial order or model family.

Details

This class is virtual and cannot be instantiated directly; it exists so that S4 methods (e.g. `show`) can be defined once and inherited by every concrete subclass.

Slots

`coefficients` numeric vector of estimated parameters
`residuals` numeric vector of final residuals/innovations
`convergence` logical or integer indicating algorithm convergence
`iterations` numeric number of iterations performed
`call` original function call

PMMtsfit-class

Virtual S4 class for PMM time-series fit objects

Description

Common parent of every concrete time-series fit class (TS2fit, TS3fit and their AR/MA/ARMA/ARIMA/seasonal subclasses). Provides the time-series-specific slots so that methods such as `predict`, `plot`, and `forecast` can be defined once and inherited.

Details

This class is virtual and inherits from [PMMfit](#).

Slots

model_type character string identifying the model family (one of "ar", "ma", "arma", "arima", "sar", "sma", "sarma", "sarima")

intercept numeric scalar intercept/mean term

original_series numeric vector of the original time series

order list of order parameters (entries depend on model_type; see the relevant fitting function)

pmm_ar

*Fit an AR model with the polynomial maximization method***Description**

Unified entry point for PMM2 and PMM3 autoregressive estimation.

Usage

```
pmm_ar(x, order, method = c("pmm2", "pmm3"), ...)
```

Arguments

x univariate time series or numeric vector.

order non-negative integer giving the autoregressive order.

method One of "pmm2" or "pmm3".

... Additional arguments forwarded to [ar_pmm2](#) or [ar_pmm3](#).

Value

An S4 object of class [ARPM2](#) (for method = "pmm2") or [ARPM3](#).

pmm_arima

*Fit an ARIMA model with the polynomial maximization method***Description**

Unified entry point for PMM2 and PMM3 ARIMA estimation. This is the recommended primary API for non-Gaussian ARIMA modelling and the interface featured in the JSS submission accompanying the package.

Usage

```
pmm_arima(x, order, method = c("pmm2", "pmm3"), ...)
```

Arguments

x	univariate time series or numeric vector.
order	integer vector c(p, d, q) giving the AR order, degree of differencing, and MA order.
method	One of "pmm2" or "pmm3".
...	Additional arguments forwarded to arma_pmm2 or arma_pmm3 .

Value

An S4 object of class [ARIMAPMM2](#) or [ARIMAPMM3](#).

Examples

```
set.seed(2026)
x <- arima.sim(list(ar = 0.6), n = 200,
               rand.gen = function(k) rgamma(k, 2, 1) - 2)
pmm_arma(x, order = c(1, 0, 0), method = "pmm2")
```

pmm_arma

Fit an ARMA model with the polynomial maximization method

Description

Unified entry point for PMM2 and PMM3 ARMA estimation.

Usage

```
pmm_arma(x, order, method = c("pmm2", "pmm3"), ...)
```

Arguments

x	univariate time series or numeric vector.
order	integer vector c(p, q) giving the AR and MA orders.
method	One of "pmm2" or "pmm3".
...	Additional arguments forwarded to arma_pmm2 or arma_pmm3 .

Value

An S4 object of class [ARMAPMM2](#) or [ARMAPMM3](#).

pmm_dispatch	<i>Automatic PMM method selection</i>
--------------	---------------------------------------

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

An object of S3 class "PMMdispatch": a list with components accessible via \$ and a dedicated print method. 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 $\gamma_6 = m_6/m_2^3 - 15m_4/m_2^2 + 30$ 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_lm

*Fit a linear model with the polynomial maximization method***Description**

Unified entry point for PMM2 and PMM3 linear regression. Dispatches to [lm_pmm2](#) or [lm_pmm3](#) based on the method argument. With method = "auto", the [pmm_dispatch](#) rule is applied to the residuals of an initial OLS fit to select the recommended method.

Usage

```
pmm_lm(formula, data, method = c("auto", "pmm2", "pmm3"), ...)
```

Arguments

formula	R formula for the model.
data	data.frame containing the variables in formula.
method	One of "auto", "pmm2", or "pmm3". "auto" consults pmm_dispatch and falls back to PMM2 when the dispatcher recommends OLS, so the return value is always a PMMfit object.
...	Additional arguments forwarded to the underlying fitter (lm_pmm2 or lm_pmm3).

Value

An S4 object of class [PMM2fit](#) or [PMM3fit](#), both inheriting from [PMMfit](#).

See Also

[pmm_dispatch](#), [pmm_arima](#), [lm_pmm2](#), [lm_pmm3](#).

Examples

```
set.seed(123)
n <- 80
x <- rnorm(n)
y <- 2 + 3 * x + rgamma(n, 2, 1) - 2
dat <- data.frame(y = y, x = x)
fit <- pmm_lm(y ~ x, data = dat, method = "pmm2")
fit
```

pmm_ma	<i>Fit an MA model with the polynomial maximization method</i>
--------	--

Description

Unified entry point for PMM2 and PMM3 moving-average estimation.

Usage

```
pmm_ma(x, order, method = c("pmm2", "pmm3"), ...)
```

Arguments

x	univariate time series or numeric vector.
order	non-negative integer giving the moving-average order.
method	One of "pmm2" or "pmm3".
...	Additional arguments forwarded to ma_pmm2 or ma_pmm3 .

Value

An S4 object of class [MAPMM2](#) or [MAPMM3](#).

pmm_sarima	<i>Fit a seasonal ARIMA model with the polynomial maximization method</i>
------------	---

Description

Unified entry point for SARIMA-PMM2 estimation. PMM3 seasonal estimation is not yet implemented; method = "pmm3" therefore raises an error. All arguments other than method are forwarded verbatim to [sarima_pmm2](#) – see that function for the seasonal order specification.

Usage

```
pmm_sarima(x, ..., method = c("pmm2", "pmm3"))
```

Arguments

x	univariate time series or numeric vector.
...	Arguments forwarded to sarima_pmm2 (notably order and seasonal).
method	Currently only "pmm2" is supported.

Value

An S4 object of class [SARIMAPMM2](#).

pmm_skewness	<i>Calculate skewness from data</i>
--------------	-------------------------------------

Description

Calculate skewness from data

Usage

```
pmm_skewness(x)
```

Arguments

x	numeric vector
---	----------------

Value

Skewness value

predict,PMM2fit-method	<i>Prediction method for PMM2fit objects</i>
------------------------	--

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 $X \%*\% \text{beta}$ where X is the design matrix constructed from newdata using the model formula, and beta 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

<code>print.PMMdispatch</code>	<i>Print method for PMMdispatch objects</i>
--------------------------------	---

Description

Print method for PMMdispatch objects

Usage

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

Arguments

<code>x</code>	object of class PMMdispatch returned by pmm_dispatch
<code>...</code>	additional arguments (ignored)

Value

The object x, invisibly.

<code>residuals,PMM2fit-method</code>	<i>Extract residuals from PMM2fit object</i>
---------------------------------------	--

Description

Extract residuals from PMM2fit object

Usage

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

Arguments

<code>object</code>	PMM2fit object
<code>...</code>	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

<code>object</code>	PMM3fit object
<code>...</code>	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

<code>object</code>	TS2fit object
<code>...</code>	Additional arguments (not used)

Value

Vector of residuals (innovations)

residuals,TS3fit-method
<i>Extract residuals from TS3fit object</i>

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	<i>S4 class for Seasonal ARIMA model results with PMM2</i>
------------------	--

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:

$$(1-\phi_1B-\dots-\phi_pB^p)(1-\Phi_1B^s-\dots-\Phi_PB^{Ps})(1-B)^d(1-B^s)^Dy_t = (1+\theta_1B+\dots+\theta_qB^q)(1+\Theta_1B^s+\dots+\Theta_QB^{Qs})\epsilon_t.$$

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: • 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) • 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

[sarima_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: • 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: • 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

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

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")
```

show,PMM2fit-method *Show method for PMM2fit objects*

Description

Prints a concise lm-style display: the original call, coefficients, and a one-line algorithm footer. Use [summary](#) for full diagnostics including moments, the g2 efficiency factor, and bootstrap inference.

Usage

```
## S4 method for signature 'PMM2fit'
show(object)
```

Arguments

object object of class PMM2fit

Value

The object (invisibly).

show,PMM3fit-method *Show method for PMM3fit objects*

Description

Prints a concise lm-style display: the original call, coefficients, and a one-line algorithm footer. Use [summary](#) for full diagnostics including moments, gamma4/gamma6 cumulants, and the g3 efficiency factor.

Usage

```
## S4 method for signature 'PMM3fit'
show(object)
```

Arguments

object object of class PMM3fit

Value

The object (invisibly).

show,TS2fit-method	Show method for TS2fit objects (and subclasses)
--------------------	---

Description

Inherited by ARPMM2, MAPMM2, ARMAPMM2, ARIMAPMM2 and the seasonal SARPMM2/SMAPMM2/SARMAPMM2 subclasses. Use [summary](#) for full diagnostics.

Usage

```
## S4 method for signature 'TS2fit'
show(object)
```

Arguments

object object inheriting from TS2fit

Value

The object (invisibly).

show,TS3fit-method	Show method for TS3fit objects (and subclasses)
--------------------	---

Description

Inherited by ARPMM3, MAPMM3, ARMAPMM3, ARIMAPMM3. Use [summary](#) for full diagnostics.

Usage

```
## S4 method for signature 'TS3fit'
show(object)
```

Arguments

object object inheriting from TS3fit

Value

The object (invisibly).

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

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

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

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

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 $J_{ij} = d(y_hat_i)/d(\theta_j)$. 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

<code>object</code>	object of class "PMM2fit"
<code>formula</code>	(optional) formula used for the model
<code>data</code>	(optional) data used
<code>B</code>	number of bootstrap replications for statistical inference
<code>...</code>	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

<code>object</code>	PMM3fit object
<code>...</code>	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

<code>object</code>	object of class "SARIMAPMM2"
<code>...</code>	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

<code>object</code>	object of class "SARMAPMM2"
<code>...</code>	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)

summary.PMMdispatch	<i>Summary method for PMMdispatch objects</i>
---------------------	---

Description

Returns a one-row data frame suitable for inclusion in reports or comparison tables.

Usage

```
## S3 method for class 'PMMdispatch'
summary(object, ...)
```

Arguments

object	object of class PMMdispatch
...	additional arguments (ignored)

Value

A data frame with columns method, n, gamma3, gamma4, gamma6, g2, g3, g_selected, improvement_pct.

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

Description

Computes the skewness coefficient gamma3 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 gamma3 (default 0.3)

Value

A list with components:

gamma3	Sample skewness coefficient
is_symmetric	Logical: TRUE if gamma3 <= threshold
message	Human-readable verdict

TS2fit-class

S4 class for PMM2 time-series fit results

Description

Common parent of every PMM2 time-series fit class (AR, MA, ARMA, ARIMA, and the seasonal SAR/SMA/SARMA/SARIMA subclasses). Inherits the PMM2-specific central moments from [BasePMM2](#) and the time-series slots (model_type, intercept, original_series, order) from [PMMtsfit](#). The diamond inheritance resolves at [PMMfit](#), which both branches share as their virtual root.

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

Slots

ma_solver character, which treatment of the moving-average recursion produced the fit: "linearized" (design matrix frozen at the CSS residuals) or "recursive" (innovation recursion recomputed at every candidate parameter). Pure AR fits are unaffected by the distinction and carry the default "linearized".

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

S4 class for PMM3 time-series fit results

Description

Common parent of every PMM3 time-series fit class (ARPM3, MAPMM3, ARMAPMM3, ARIMAPMM3). Inherits the PMM3-specific moments and cumulant coefficients from [BasePMM3](#) and the time-series slots from [PMMtsfit](#). The diamond inheritance resolves at [PMMfit](#), which both branches share as their virtual root.

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",
  ma_solver = c("linearized", "recursive"),
  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"
ma_solver	String: how the moving-average recursion is treated, "linearized" (default) or "recursive". Has no effect when the MA order is zero. See Details.
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

Treatment of the MA recursion (ma_solver).

- "linearized" (default, historical behaviour): the design matrix is built once from the CSS residuals and held fixed. For AR columns this is exact, because the regressors are observed data. For MA columns it is a one-step linearisation of the innovation recursion and costs a factor of exactly $1 - \theta^2$ in asymptotic variance relative to CSS. The penalty does *not* vanish as n grows.
- "recursive" (recommended whenever $q > 0$): the innovations $\varepsilon_t(\beta)$ and the score regressors $x_t(\beta) = -\partial \varepsilon_t / \partial \beta$ are recomputed by the exact recursion at every candidate parameter. When $\mu_3 = 0$ the estimating equation then reduces identically to the CSS first-order condition, so the $1 - \theta^2$ penalty disappears exactly rather than asymptotically. Measured on ARIMA(0,1,1) with $\theta = -0.5$, $T = 500$: relative efficiency against CSS rises from 0.77 to 0.98 (Gaussian), 1.30 to 1.72 (Gamma), 1.33 to 1.81 (lognormal) and 1.35 to 1.73 (chi-squared).

The default is "linearized" for backward compatibility only; it will change in a future major release. `ma_solver` is supported for `model_type` "ma", "arma" and "arima" with $q \geq 1$ and any $p \geq 0$. Seasonal models ([sarima_pmm2](#) and friends) always use the linearised path.

Not to be confused with the unrelated `ma_method` argument of [sarima_pmm2](#), which takes "mle"/"pmm2" and selects how the seasonal MA terms are estimated. The two control different things and are not interchangeable.

Value

An S4 object `TS2fit` of the corresponding subclass

ts_pmm2_inference

Bootstrap inference for PMM2 time series models

Description

Residual- or block-bootstrap standard errors and confidence intervals for PMM2 time-series fits. Block bootstrap can exhibit substantial finite-sample bias for AR coefficients near the stationarity boundary; see `ci_method` and the returned `bias` column for diagnostics.

Usage

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

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
ci_method	character: confidence-interval method. See pmm2_inference for the full description. The default "normal" returns a Wald interval centred on the point estimate, which is robust to the moderate downward bias that block bootstrap induces for AR coefficients of strongly persistent series. Use "percentile" for Efron's classical percentile interval or "basic" for the Davison & Hinkley (1997) pivotal interval.

Value

data.frame with columns: Estimate, Std.Error, bias, t.value, p.value, conf.low, conf.high.

ts_pmm3	<i>Fit a time series model using PMM3</i>
---------	---

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 ts_pmm2 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 κ from initial residuals
3. Checks symmetry: warns if $|\text{gamma}_3| > 0.3$
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

vcov,PMM2fit-method	<i>Variance-covariance matrix for PMM2fit object</i>
---------------------	--

Description

Returns the asymptotic covariance matrix $\sigma^2 g_2 (X^\top X)^{-1}$, where $g_2 = 1 - c_3^2 / (2 + c_4)$ is the PMM2 variance reduction factor. Under Gaussian errors $g_2 = 1$ and the result equals the OLS covariance matrix.

Usage

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

Arguments

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

Value

Numeric matrix of the same dimension as the number of coefficients

vcov,PMM3fit-method	<i>Variance-covariance matrix for PMM3fit objects</i>
---------------------	---

Description

Returns the asymptotic covariance matrix $V_{\text{PMM3}} = g_3 \sigma^2 (X^\top X)^{-1}$ where $g_3 = 1 - \gamma_4^2 / (6 + 9\gamma_4 + \gamma_6)$ is the PMM3 variance reduction factor (Kunchenko's polynomial maximization method, $s = 3$ specialised to symmetric platykurtic errors; see [pmm3_variance_matrices](#)). Under Gaussian errors $g_3 = 1$ and the result equals the OLS covariance.

Usage

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

Arguments

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

Value

Numeric matrix of the same dimension as the number of coefficients

vcov,TS2fit-method	Variance-covariance matrix for TS2fit models
--------------------	--

Description

For AR models, returns $\sigma^2 g_2(X^\top X)^{-1}$ where X is the lagged design matrix.

Usage

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

Arguments

object	TS2fit (or subclass) object
...	Additional arguments (not used)

Details

For non-seasonal MA/ARMA/ARIMA models fitted with `ma_solver = "recursive"` the same expression applies with X replaced by the exact score regressors $x_t = -\partial \varepsilon_t / \partial \beta$, rebuilt by `pmm2_recursive_design()`. This is the closed-form sandwich $\Sigma = (\Delta/a) \mathbf{M}^{-1}$ with $a = \mu_4 - \mu_2^2$, $\Delta = \mu_2 a - \mu_3^2$ and $\mathbf{M} = E[x_t x_t^\top]$; the estimating function is a martingale difference sequence at the true parameter, so no long-run (HAC) correction is required and no numerical Jacobian is needed. Note $\Delta/a = \mu_2 g_2$, i.e. the AR and MA branches share one formula.

For fits produced with the (default) `ma_solver = "linearized"` the frozen CSS design matrix is not the derivative of the criterion, the sandwich above does not apply, and `vcov()` still refuses; use [ts_pmm2_inference](#) for bootstrap standard errors.

Value

Numeric covariance matrix of the slope coefficients

Known limitation (intercept)

Neyman orthogonality with respect to the plug-in moments holds if and only if $\mu_3 E[x_t] = 0$. All slope regressors have $E[x_t] = 0$, so estimating $\hat{\mu}_2, \hat{\mu}_3, \hat{\mu}_4$ does not affect the limiting distribution of the AR/MA coefficients. The intercept regressor has $E[x_t] = 1$, so under skewed innovations ($\mu_3 \neq 0$) estimating $\hat{\mu}_2$ contributes at first order to the intercept and its reported standard error is too small. This affects the linearised branch as well. `vcov()` therefore covers the slope parameters only and warns when an intercept was estimated under visible skewness.

vcov,TS3fit-method	<i>Variance-covariance matrix for TS3fit AR models</i>
--------------------	--

Description

For AR models, returns $\sigma^2 g_3 (X^\top X)^{-1}$ where X is the lagged design matrix and $g_3 = 1 - \gamma_4^2 / (6 + 9\gamma_4 + \gamma_6)$ is the PMM3 efficiency factor (Kunchenko, s = 3 specialised to symmetric platykurtic innovations). For MA/ARMA/ARIMA models the asymptotic covariance requires the full Fisher information matrix; use bootstrap standard errors via [ts_pmm2_inference](#) on an analogous PMM2 fit, or refit with [ar_pmm3](#) on the residual series.

Usage

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

Arguments

object	TS3fit (or subclass) object
...	Additional arguments (not used)

Value

Numeric covariance matrix (AR models only)

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