

Package ‘EmpiricalDynamics’

January 16, 2026

Title Empirical Discovery of Differential Equations from Time Series Data

Version 0.1.2

Description A comprehensive toolkit for discovering differential and difference equations from empirical time series data using symbolic regression. The package implements a complete workflow from data preprocessing (including Total Variation Regularized differentiation for noisy economic data), visual exploration of dynamical structure, and symbolic equation discovery via genetic algorithms. It leverages a high-performance 'Julia' backend ('SymbolicRegression.jl') to provide industrial-grade robustness, physics-informed constraints, and rigorous out-of-sample validation. Designed for economists, physicists, and researchers studying dynamical systems from observational data.

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Encoding UTF-8

RoxygenNote 7.3.3

SystemRequirements Julia (>= 1.6)

Depends R (>= 4.0.0)

Imports JuliaCall (>= 0.17), CVXR (>= 1.0), minpack.lm (>= 1.2), signal (>= 0.7), lmtest (>= 0.9), tseries (>= 0.10), ggplot2 (>= 3.4.0), gridExtra (>= 2.3), stats, graphics, grDevices, utils, methods

Suggests osqp (>= 0.6), ECOSolveR (>= 0.5), testthat (>= 3.0.0), knitr (>= 1.40), rmarkdown (>= 2.20), covr, mgcv

Config/testthat.edition 3

VignetteBuilder knitr

URL <https://github.com/IsadoreNabi/EmpiricalDynamics>

BugReports <https://github.com/IsadoreNabi/EmpiricalDynamics/issues>

NeedsCompilation no

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Repository CRAN

Date/Publication 2026-01-16 11:30:34 UTC

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Description

Creates a concise text summary of the analysis.

Usage

```
analysis_summary(results, verbose = TRUE)
```

Arguments

- | | |
|---------|-----------------------------|
| results | Analysis results list. |
| verbose | Include additional details. |

Value

Character string with summary.

analyze_bifurcations *Analyze Bifurcations*

Description

Examines how fixed points change as a parameter varies.

Usage

```
analyze_bifurcations(
  equation,
  variable,
  parameter,
  param_range = c(-5, 5),
  n_param = 50,
  z_range = c(-10, 10),
  exogenous_values = list()
)
```

Arguments

- | | |
|------------------|-------------------------------------|
| equation | Fitted equation object. |
| variable | Name of the main variable. |
| parameter | Name of the parameter to vary. |
| param_range | Range for parameter values. |
| n_param | Number of parameter values to test. |
| z_range | Range for searching fixed points. |
| exogenous_values | Fixed values for other variables. |

Value

Object of class "bifurcation_analysis".

 analyze_fixed_points *Analyze Fixed Points*

Description

Finds and characterizes fixed points of the discovered equation.

Usage

```
analyze_fixed_points(
  equation,
  variable,
  range = c(-10, 10),
  n_grid = 100,
  exogenous_values = list()
)
```

Arguments

equation	Fitted equation object.
variable	Name of the main variable.
range	Numeric vector of length 2 specifying search range.
n_grid	Number of grid points for initial search.
exogenous_values	Named list of fixed values for exogenous variables.

Value

Data frame of fixed points with stability classification.

Examples

```
# Toy example: dZ = 2*Z - Z^2 (Logistic growth)
data <- data.frame(Z = seq(0.1, 3, length.out=50))
data$dZ <- 2*data$Z - data$Z^2
model <- stats::lm(dZ ~ I(Z) + I(Z^2) + 0, data = data)

# Analyze (note: linear models on dZ aren't direct ODEs, but this demonstrates structure)
# For correct usage, 'equation' should be from fit_specified_equation
fp <- analyze_fixed_points(model, variable = "Z", range = c(0, 3))
```

`annotate_hypotheses` *Annotate Hypotheses*

Description

Records researcher hypotheses based on visual exploration, to be used as constraints or guides in subsequent symbolic search.

Usage

```
annotate_hypotheses(data, hypotheses)
```

Arguments

<code>data</code>	Data frame (with exploration results as attribute).
<code>hypotheses</code>	Character vector of hypotheses.

Value

Data frame with hypotheses attached as attribute.

Examples

```
# Toy example
data <- data.frame(Z = 1:10)
data <- annotate_hypotheses(data, c(
  "Z exhibits logistic saturation around Z=100",
  "Effect of X appears linear"
))
```

`bootstrap_parameters` *Bootstrap Confidence Intervals for Parameters*

Description

Computes bootstrap confidence intervals for equation parameters.

Usage

```
bootstrap_parameters(  
  equation,  
  data,  
  response = NULL,  
  derivative_col = NULL,  
  n_boot = 500,  
  conf_level = 0.95,  
  block_size = NULL  
)
```

Arguments

equation	Fitted equation object.
data	Original data.
response	Name of response column.
derivative_col	Alias for response.
n_boot	Number of bootstrap samples.
conf_level	Confidence level (default 0.95).
block_size	Block size for block bootstrap (time series).

Value

Data frame with parameter estimates and confidence intervals.

check_qualitative_behavior

Check Qualitative Behavior

Description

Comprehensive check of whether the discovered equation exhibits expected qualitative features.

Usage

```
check_qualitative_behavior(  
  equation,  
  data,  
  variable,  
  expected_features = list()  
)
```

Arguments

- equation Fitted equation object.
 data Original data.
 variable Main variable name.
 expected_features
 List of expected qualitative features:
 • n_fixed_points: Expected number of fixed points
 • stability_pattern: e.g., c("stable", "unstable", "stable")
 • monotonicity: Expected sign of derivative ("positive", "negative", "none")
 • bounded: Whether dynamics should be bounded

Value

Object of class "qualitative_check".

coefficient_table *Generate Coefficient Table*

Description

Creates a publication-ready table of estimated coefficients.

Usage

```
coefficient_table(  

  equation,  

  bootstrap_results = NULL,  

  format = c("data.frame", "latex", "markdown", "html"),  

  caption = "Estimated Coefficients",  

  label = "tab:coefficients"  

)
```

Arguments

- equation Fitted equation object.
 bootstrap_results
 Results from bootstrap_parameters (optional).
 format Output format: "data.frame", "latex", "markdown", "html".
 caption Table caption.
 label LaTeX label for referencing.

Value

Formatted table.

```
compare_differentiation_methods
    Compare Differentiation Methods
```

Description

Applies multiple differentiation methods to the same data and produces a comparison plot.

Usage

```
compare_differentiation_methods(
  Z,
  t = NULL,
  methods = c("tvr", "savgol", "spline", "finite_diff"),
  plot = TRUE
)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.
methods	Character vector of methods to compare.
plot	Produce comparison plot?

Value

List of derivative vectors from each method.

```
compare_estimation_methods
    Compare OLS and GLS Estimation
```

Description

Produces a comparison of drift estimates from ordinary least squares versus iterative GLS.

Usage

```
compare_estimation_methods(ols_model, gls_model, data)
```

Arguments

ols_model	SDE model estimated with OLS
gls_model	SDE model estimated with iterative GLS
data	Data frame used for estimation

Value

Invisibly returns comparison statistics

compare_trajectories *Compare Simulated and Observed Trajectories*

Description

Computes metrics comparing simulated trajectories to observed data.

Usage

```
compare_trajectories(
  simulation,
  observed_data,
  time_col = "time",
  var_col = NULL
)
```

Arguments

<code>simulation</code>	Trajectory simulation object.
<code>observed_data</code>	Data frame with observed values.
<code>time_col</code>	Name of time column.
<code>var_col</code>	Name of variable column to compare.

Value

Data frame with comparison metrics.

compute_derivative *Compute Derivative of a Time Series*

Description

Main dispatcher function for numerical differentiation. Supports multiple methods appropriate for different data characteristics.

Usage

```
compute_derivative(
  Z,
  t = NULL,
  method = c("tvr", "savgol", "spline", "finite_diff", "spectral"),
  ...
)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points (NULL assumes dt=1).
method	Differentiation method: "tvr", "savgol", "spline", "finite_diff", "fd", or "spectral".
...	Additional arguments passed to the specific method.

Details

Available methods:

- **tvr**: Total Variation Regularized differentiation (recommended for economic data with trends and shocks).
- **savgol**: Savitzky-Golay filter (moderate noise, preserves peaks).
- **spline**: Smoothing spline (high noise, prioritizes trend).
- **finite_diff** or **fd**: Centered finite differences (low noise).
- **spectral**: FFT-based (periodic data only).

Value

Numeric vector of estimated derivatives with diagnostic attributes.

See Also

[compute_derivative_tvr](#), [suggest_differentiation_method](#)

Examples

```
t <- seq(0, 10, by = 0.1)
Z <- sin(t) + rnorm(length(t), sd = 0.1)

# Finite differences (fast, no dependencies)
dZ_fd <- compute_derivative(Z, t, method = "finite_diff")

# Access the derivative vector for plotting
plot(t, dZ_fd$derivative, type = "l", main = "Derivative Comparison")
lines(t, cos(t), col = "red", lty = 2) # True derivative

# TVR (requires CVXR)
if (requireNamespace("CVXR", quietly = TRUE)) {
  dZ_tvr <- compute_derivative(Z, t, method = "tvr")
}
```

compute_derivatives *Compute Derivatives for Specified Variables*

Description

Convenience function to compute derivatives for all endogenous variables in a specified dataset.

Usage

```
compute_derivatives(data, method = "tvr", prefix = "d_", ...)
```

Arguments

<code>data</code>	Data frame with variable specifications (from <code>specify_variables</code>).
<code>method</code>	Differentiation method.
<code>prefix</code>	Prefix for derivative column names (default "d_").
<code>...</code>	Additional arguments passed to <code>compute_derivative</code> .

Value

Data frame with derivative columns added.

compute_derivative_fd *Centered Finite Differences*

Description

Computes derivatives using centered finite differences.

Usage

```
compute_derivative_fd(Z, t = NULL, ...)
```

Arguments

<code>Z</code>	Numeric vector of observations.
<code>t</code>	Numeric vector of time points.
<code>...</code>	Additional arguments (ignored).

Value

List with derivative vector.

```
compute_derivative_savgol
    Savitzky-Golay Derivative
```

Description

Computes derivatives using the Savitzky-Golay filter.

Usage

```
compute_derivative_savgol(Z, t = NULL, p = 3, n = NULL, m = 1, ...)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.
p	Polynomial order (default 3).
n	Filter length (must be odd, default auto-selected).
m	Derivative order (default 1).
...	Additional arguments (ignored).

Value

List with derivative vector.

```
compute_derivative_spectral
    Spectral (FFT) Differentiation
```

Description

Computes derivatives using the Fourier transform.

Usage

```
compute_derivative_spectral(Z, t = NULL, ...)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.
...	Additional arguments (ignored).

Value

List with derivative vector.

Warning

This method assumes the signal is periodic.

compute_derivative_spline

Smoothing Spline Derivative

Description

Computes derivatives by fitting a smoothing spline and differentiating it.

Usage

```
compute_derivative_spline(Z, t = NULL, spar = NULL, df = NULL, ...)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.
spar	Smoothing parameter (NULL for automatic selection).
df	Degrees of freedom (alternative to spar).
...	Additional arguments (ignored).

Value

List with derivative vector.

compute_derivative_tvr

Total Variation Regularized Differentiation

Description

Computes derivatives by solving a convex optimization problem that balances fidelity to the data against smoothness of the derivative.

Usage

```
compute_derivative_tvr(Z, t = NULL, lambda = "auto", solver = "osqp", ...)
```

Arguments

<code>z</code>	Numeric vector of observations.
<code>t</code>	Numeric vector of time points (NULL assumes dt=1).
<code>lambda</code>	Regularization parameter ("auto" for cross-validation selection).
<code>solver</code>	Optimization backend: "osqp", "ecos", or "scs".
<code>...</code>	Additional arguments (ignored).

Value

Object of class "tvr_derivative" (also a list with \$derivative).

`compute_residuals` *Compute Residuals from Symbolic Equation*

Description

Calculates residuals from a fitted symbolic equation or SDE model.

Usage

```
compute_residuals(model, data, target = NULL)
```

Arguments

<code>model</code>	A <code>symbolic_equation</code> or <code>sde_model</code> object
<code>data</code>	Data frame containing the variables
<code>target</code>	Name of target variable (auto-detected if NULL)

Value

Numeric vector of residuals

construct_sde*Construct Stochastic Differential Equation Model*

Description

Combines a drift equation and diffusion model into a complete SDE: $dZ = f(Z, X, Y) dt + g(Z, X, Y) dW$

Usage

```
construct_sde(
  drift,
  diffusion = NULL,
  variable = NULL,
  refine_with_gls = FALSE,
  gls_max_iter = 10,
  gls_tolerance = 1e-04,
  data = NULL,
  target = NULL
)
```

Arguments

<code>drift</code>	Symbolic equation for the drift term $f(\cdot)$
<code>diffusion</code>	Variance model for the diffusion term $g(\cdot)$
<code>variable</code>	Name of the main state variable
<code>refine_with_gls</code>	Use iterative GLS to refine estimates?
<code>gls_max_iter</code>	Maximum iterations for GLS
<code>gls_tolerance</code>	Convergence tolerance for GLS
<code>data</code>	Data frame (required if <code>refine_with_gls = TRUE</code>)
<code>target</code>	Target variable name (required if <code>refine_with_gls = TRUE</code>)

Value

An object of class "sde_model"

create_transformations

Create Candidate Transformations

Description

Generates derived variables based on specified transformations that may be theoretically relevant.

Usage

```
create_transformations(  
  data,  
  transformations = NULL,  
  variables = NULL,  
  cols = NULL  
)
```

Arguments

data	Data frame.
transformations	List of formulas specifying transformations.
variables	Character vector of variable names (alternative interface).
cols	Alias for variables.

Value

Data frame with transformation columns added.

Examples

```
data <- data.frame(X = 1:10, Y = 10:1)  
  
# Simple interface  
data <- create_transformations(data, variables = c("X", "Y"))  
  
# Formula interface  
data <- create_transformations(data, transformations = list(  
  ratios = ~ X/Y  

```

cross_validate	<i>Cross-Validate Discovered Equation</i>
----------------	---

Description

Performs k-fold or block cross-validation to assess out-of-sample predictive performance of the discovered equation.

Usage

```
cross_validate(
  equation,
  data,
  response = NULL,
  derivative_col = NULL,
  k = 5,
  method = c("block", "random", "rolling"),
  block_size = NULL,
  horizon = 1,
  refit_derivative = FALSE,
  diff_method = "tvr",
  verbose = TRUE
)
```

Arguments

<code>equation</code>	Fitted equation object from <code>fit_specified_equation</code> or <code>symbolic_search</code> . Can also be an object of class <code>lm</code> or <code>nls</code> .
<code>data</code>	Data frame containing all variables.
<code>response</code>	Name of the response column (derivative).
<code>derivative_col</code>	Alias for <code>response</code> (for compatibility).
<code>k</code>	Number of folds for cross-validation.
<code>method</code>	CV method: "random", "block", "rolling".
<code>block_size</code>	For block methods, size of contiguous blocks.
<code>horizon</code>	For rolling CV, forecast horizon.
<code>refit_derivative</code>	Logical; whether to recompute derivatives for each fold (currently unused).
<code>diff_method</code>	Differentiation method if refitting (currently unused).
<code>verbose</code>	Print progress.

Value

Object of class "cv_result" containing:

rmse	Root mean squared error per fold
mae	Mean absolute error per fold
r_squared	R-squared per fold
mean_rmse	Average RMSE across folds
sd_rmse	Standard deviation of RMSE
predictions	List of predicted vs actual per fold
fold_indices	Indices used for each fold

Examples

```
# Toy example using lm
data <- data.frame(
  time = 1:50,
  y = seq(1, 10, length.out = 50) + stats::rnorm(50, sd = 0.1)
)
# Simple linear model as a proxy for a discovered equation
model <- stats::lm(y ~ time, data = data)

# Run cross-validation
cv_res <- cross_validate(
  equation = model,
  data = data,
  response = "y",
  k = 3,
  method = "random"
)
print(cv_res)
```

define_custom_operators

Define Custom Operators

Description

Defines custom mathematical operators for use in symbolic search.

Usage

`define_custom_operators(...)`

Arguments

... Named functions to add as operators.

Value

List of operator definitions suitable for symbolic_search.

Examples

```
ops <- define_custom_operators(
  logistic = function(x, k = 1, x0 = 0) 1 / (1 + exp(-k * (x - x0))),
  threshold = function(x, c) ifelse(x > c, 1, 0)
)
```

diagnose_sampling_frequency

Diagnose Sampling Frequency

Description

Evaluates whether the sampling frequency is appropriate for capturing the dynamics of the phenomenon.

Usage

```
diagnose_sampling_frequency(Z, t = NULL)
```

Arguments

- | | |
|---|---------------------------------|
| Z | Numeric vector of observations. |
| t | Numeric vector of time points. |

Value

List with diagnostic information and recommendations.

ed_theme

Default ggplot2 Theme for EmpiricalDynamics

Description

A clean, publication-ready theme for all diagnostic plots.

Usage

```
ed_theme(base_size = 11, base_family = "")
```

Arguments

- base_size Base font size.
base_family Base font family.

Value

A ggplot2 theme object.

estimate_initial_values

Automatic Initial Value Estimation

Description

Estimates reasonable starting values for nonlinear least squares.

Usage

```
estimate_initial_values(  
  expression,  
  data,  
  response = NULL,  
  derivative_col = NULL,  
  method = c("grid_search", "random", "heuristic"),  
  n_tries = 100  
)
```

Arguments

- expression Character string with the equation expression.
data Data frame with variables.
response Name of the response variable.
derivative_col Alias for response (for compatibility).
method Estimation method: "grid_search", "random", or "heuristic".
n_tries Number of attempts for random method.

Value

Named list of initial values.

`estimate_sde_iterative`

Iterative GLS Estimation for SDEs

Description

Refines drift and diffusion estimates using iterative Generalized Least Squares, which is more appropriate when heteroscedasticity is substantial.

Usage

```
estimate_sde_iterative(
  target,
  predictors,
  data,
  initial_drift = NULL,
  max_iter = 10,
  tol = 1e-04
)
```

Arguments

<code>target</code>	Numeric vector of target values (derivatives)
<code>predictors</code>	Data frame of predictor variables
<code>data</code>	Full data frame
<code>initial_drift</code>	Initial drift equation (optional)
<code>max_iter</code>	Maximum number of iterations
<code>tol</code>	Convergence tolerance (RMSE change in coefficients)

Value

An `sde_model` object with refined estimates

`exploration`

Visual Exploration of Dynamical Structure

Description

Functions for visually exploring the structure of dynamical systems before formal model fitting. These diagnostics should be used BEFORE any symbolic search to inform hypotheses about functional forms.

<code>explore_dynamics</code>	<i>Comprehensive Dynamics Exploration</i>
-------------------------------	---

Description

Generates a battery of diagnostic plots to explore the dynamical structure of the data and suggests potential functional forms.

Usage

```
explore_dynamics(
  data,
  target,
  predictors = NULL,
  time = NULL,
  n_bins = 10,
  include = "all"
)
```

Arguments

<code>data</code>	Data frame containing the time series.
<code>target</code>	Name of the target variable (or its derivative).
<code>predictors</code>	Character vector of predictor variable names.
<code>time</code>	Name of the time column (auto-detected if NULL).
<code>n_bins</code>	Number of bins for conditional analysis.
<code>include</code>	Which plots to include: "all", or subset of c("timeseries", "phase", "bivariate", "interactions").

Value

A list containing:

- `suggestions`: Character vector of suggested functional forms
- `statistics`: Data frame of diagnostic statistics
- `plots`: List of ggplot objects (if available)

Examples

```
# Toy example
data <- data.frame(
  time = 1:50,
  Z = sin(seq(0, 10, length.out = 50)),
  X = cos(seq(0, 10, length.out = 50))
)
data$dZ <- c(diff(data$Z)/diff(data$time), NA)
```

```
data <- na.omit(data)

result <- explore_dynamics(data,
  target = "dZ",
  predictors = c("Z", "X")
)
print(result$suggestions)
```

export_results *Export Results to Multiple Formats*

Description

Exports analysis results to various file formats.

Usage

```
export_results(
  results,
  output_dir,
  prefix = "empirical_dynamics",
  formats = c("rds", "csv")
)
```

Arguments

<code>results</code>	Analysis results list.
<code>output_dir</code>	Output directory (required, no default to comply with CRAN policy).
<code>prefix</code>	File name prefix.
<code>formats</code>	Vector of formats: "rds", "csv", "json", "latex".

Value

List of paths to created files.

Examples

```
# Toy example
tmp_dir <- tempdir()
mock_results <- list(
  equation = stats::lm(mpg ~ wt, data = mtcars)
)

# Export
paths <- export_results(mock_results, output_dir = tmp_dir, formats = c("csv", "rds"))
```

```
fit_residual_distribution
    Fit Residual Distribution
```

Description

Fits candidate probability distributions to residuals, optionally with parameters that depend on state variables.

Usage

```
fit_residual_distribution(
  residuals,
  candidates = c("normal", "t", "skew-normal"),
  conditional_on = NULL,
  data = NULL
)
```

Arguments

residuals	Numeric vector of residuals
candidates	Character vector of distribution families to try
conditional_on	Formula for conditional parameters (optional)
data	Data frame (required if conditional_on specified)

Value

List with best fitting distribution and parameters

```
fit_specified_equation
    Fit Specified Equation
```

Description

Fits a researcher-specified functional form, estimating only the parameters. Uses Levenberg-Marquardt algorithm for robustness.

Usage

```
fit_specified_equation(
  expression,
  data,
  response = NULL,
  derivative_col = NULL,
  start = NULL,
  method = c("LM", "nls", "optim"),
  weights = NULL,
  lower = -Inf,
  upper = Inf
)
```

Arguments

<code>expression</code>	Character string specifying the equation (e.g., "a + b * Z").
<code>data</code>	Data frame with predictor variables.
<code>response</code>	Name of the response/target column.
<code>derivative_col</code>	Alias for response (for compatibility).
<code>start</code>	List of starting values for parameters (auto-estimated if NULL).
<code>method</code>	Optimization method: "LM" (Levenberg-Marquardt, recommended), "nls" (standard), or "optim" (general optimization).
<code>weights</code>	Optional weight vector.
<code>lower</code>	Lower bounds for parameters (for "optim" method).
<code>upper</code>	Upper bounds for parameters (for "optim" method).

Value

An object of class "symbolic_equation" containing the fitted model.

Examples

```
# Toy example
data <- data.frame(Z = seq(1, 10, length.out = 20))
data$dZ <- 0.5 * data$Z * (1 - data$Z / 20) + rnorm(20, sd = 0.01)

# Fit logistic equation
eq <- fit_specified_equation(
  expression = "r * Z * (1 - Z/K)",
  data = data,
  response = "dZ",
  start = list(r = 0.5, K = 20)
)
print(eq)
```

format_equation	<i>Format Equation for Display</i>
-----------------	------------------------------------

Description

Creates a nicely formatted string representation of the equation.

Usage

```
format_equation(  
  equation,  
  format = c("text", "latex", "markdown"),  
  precision = 4  
)
```

Arguments

equation	Fitted equation object.
format	Output format: "text", "latex", "markdown".
precision	Number of decimal places.

Value

Formatted string.

generate_report	<i>Generate Analysis Report</i>
-----------------	---------------------------------

Description

Creates a comprehensive report of the entire analysis workflow.

Usage

```
generate_report(  
  results,  
  output_file,  
  format = c("markdown", "html", "latex"),  
  title = "Empirical Dynamics Analysis Report",  
  author = "EmpiricalDynamics",  
  include_plots = TRUE  
)
```

Arguments

<code>results</code>	List containing analysis results with elements:
	<ul style="list-style-type: none"> • <code>data</code>: Original data frame • <code>derivatives</code>: Computed derivatives • <code>exploration</code>: Results from <code>explore_dynamics</code> • <code>equation</code>: Best fitted equation • <code>sde</code>: SDE model (optional) • <code>validation</code>: Results from <code>validate_model</code>
<code>output_file</code>	Path for output file (required, no default to comply with CRAN policy).
<code>format</code>	Report format: "markdown", "html", "latex".
<code>title</code>	Report title.
<code>author</code>	Author name.
<code>include_plots</code>	Include diagnostic plots.

Value

Path to generated report.

Examples

```
# Toy example to demonstrate report generation
# Using a temporary file to avoid writing to user's working directory
tmp_file <- tempfile("report_example")

# Mock results object
mock_results <- list(
  data = data.frame(time = 1:10, Z = runif(10)),
  equation = stats::lm(Z ~ time, data = data.frame(time = 1:10, Z = runif(10)))
)

# Generate report
report_path <- generate_report(mock_results, output_file = tmp_file, format = "markdown")
if(file.exists(report_path)) unlink(report_path)
```

`get_analysis_template` *Get Analysis Template*

Description

Returns a template script for running a complete analysis.

Usage

```
get_analysis_template(output_file = NULL)
```

Arguments

output_file Path to save template.

Value

Template code as character string (invisibly).

Examples

```
# Save template to a temporary file
tmp_file <- tempfile("analysis_template", fileext = ".R")
get_analysis_template(tmp_file)

# Clean up
if (file.exists(tmp_file)) unlink(tmp_file)
```

get_pareto_set *Get Full Pareto Set*

Description

Returns all equations on the Pareto front as a list.

Usage

get_pareto_set(results)

Arguments

results A symbolic_search_result object.

Value

List of symbolic_equation objects.

`list_example_data` *List Available Example Datasets*

Description

Returns information about the example datasets included with the package.

Usage

```
list_example_data()
```

Value

A data.frame with dataset names and descriptions.

Examples

```
list_example_data()
```

`load_example_data` *Load Example Dataset*

Description

Load one of the example datasets included with the package.

Usage

```
load_example_data(name)
```

Arguments

- | | |
|-------------------|--|
| <code>name</code> | Name of the dataset to load. Available datasets: <ul style="list-style-type: none">• "logistic_growth" - Logistic population growth• "predator_prey" - Lotka-Volterra predator-prey dynamics• "interest_rate" - Vasicek mean-reverting interest rate• "epidemic_data" - SIR epidemic model• "oscillator_data" - Van der Pol oscillator• "business_cycle" - Kaldor-type business cycle |
|-------------------|--|

Value

A data.frame containing the time series data.

Examples

```
# Load logistic growth data if available
if(requireNamespace("utils", quietly = TRUE)) {
  try({
    data <- load_example_data("logistic_growth")
    head(data)
  })
}
```

model_comparison_table

Generate Model Comparison Table

Description

Creates a table comparing multiple candidate equations.

Usage

```
model_comparison_table(
  equations,
  data,
  derivative_col,
  format = c("data.frame", "latex", "markdown"),
  caption = "Model Comparison"
)
```

Arguments

equations	Named list of fitted equation objects.
data	Data for computing fit statistics.
derivative_col	Response variable column.
format	Output format.
caption	Table caption.

Value

Comparison table.

`model_conditional_variance`
Model Conditional Variance

Description

Estimates how the residual variance depends on state variables, used for constructing the diffusion term of an SDE.

Usage

```
model_conditional_variance(
  residuals,
  predictors,
  data = NULL,
  method = c("symbolic", "linear", "quadratic", "gam", "constant"),
  transform = c("absolute", "squared", "log_squared"),
  ...
)
```

Arguments

<code>residuals</code>	Numeric vector of residuals
<code>predictors</code>	Formula or data frame of predictor variables (or vector of names)
<code>data</code>	Data frame (if predictors is a formula or vector of names)
<code>method</code>	Modeling method: "symbolic", "linear", "quadratic", "gam", or "constant"
<code>transform</code>	Transformation of residuals: "squared", "absolute", or "log_squared"
...	Additional arguments passed to the modeling function

Value

An object of class "variance_model" containing the fitted model

`output` *Output and Report Generation*

Description

Functions for generating publication-ready outputs including LaTeX equations, comprehensive reports, and formatted summaries.

```
plot.bifurcation_analysis  
    Plot Bifurcation Diagram
```

Description

Plot Bifurcation Diagram

Usage

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

Arguments

x Object of class bifurcation_analysis.
... Additional arguments (ignored).

Value

A ggplot object.

```
plot.cv_result      Plot CV Results
```

Description

Plot CV Results

Usage

```
## S3 method for class 'cv_result'  
plot(x, type = c("predictions", "folds", "both"), ...)
```

Arguments

x Object of class cv_result.
type Type of plot: "predictions", "folds", or "both".
... Additional arguments (ignored).

Value

A ggplot object or a list of ggplot objects.

plot.trajectory_simulation
Plot Simulated Trajectories

Description

Plot Simulated Trajectories

Usage

```
## S3 method for class 'trajectory_simulation'
plot(
  x,
  observed_data = NULL,
  show_trajectories = TRUE,
  n_show = 20,
  alpha_traj = 0.2,
  ...
)
```

Arguments

x	Object of class trajectory_simulation.
observed_data	Optional observed data to overlay.
show_trajectories	Show individual trajectories?
n_show	Number of trajectories to show.
alpha_traj	Transparency for trajectories.
...	Additional arguments (ignored).

Value

A ggplot object.

plot.tvr_derivative *Plot Method for TVR Derivative*

Description

Plot Method for TVR Derivative

Usage

```
## S3 method for class 'tvr_derivative'
plot(x, t = NULL, ...)
```

Arguments

- x A tvr_derivative object.
- t Time vector (uses attribute if NULL).
- ... Additional plot arguments.

Value

Invisibly returns the input object (called for side effects).

```
plot.validation_result  
Plot Validation Results
```

Description

Plot Validation Results

Usage

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

Arguments

- x Object of class validation_result.
- ... Additional arguments (ignored).

Value

A list of ggplot objects (invisible).

```
plot_bivariate Bivariate Scatter Plot
```

Description

Creates a scatter plot with optional nonparametric fit and marginal distributions.

Usage

```
plot_bivariate(  
  data,  
  x_var,  
  y_var,  
  color_var = NULL,  
  show_fit = TRUE,  
  show_marginals = FALSE  
)
```

Arguments

<code>data</code>	Data frame.
<code>x_var</code>	X variable name.
<code>y_var</code>	Y variable name.
<code>color_var</code>	Optional variable for color mapping.
<code>show_fit</code>	Add smooth fit?
<code>show_marginals</code>	Add marginal histograms?

Value

A ggplot object.

`plot_pareto_front` *Plot Pareto Front*

Description

Visualizes the trade-off between equation complexity and fit quality.

Usage

```
plot_pareto_front(results, highlight_selection = "knee", show_all = FALSE)
```

Arguments

<code>results</code>	A symbolic_search_result object.
<code>highlight_selection</code>	Which equation to highlight ("knee", "BIC", "AIC", or index).
<code>show_all</code>	Show all equations or just Pareto front?

Value

A ggplot object.

plot_phase_1d *1D Phase Diagram*

Description

Creates a phase diagram plotting dZ vs Z , useful for visualizing autonomous dynamics and identifying fixed points.

Usage

```
plot_phase_1d(  
  data,  
  z_var,  
  dz_var = NULL,  
  show_zero_line = TRUE,  
  show_fit = TRUE,  
  fit_method = "loess"  
)
```

Arguments

<code>data</code>	Data frame.
<code>z_var</code>	Name of state variable Z .
<code>dz_var</code>	Name of derivative dZ (auto-constructed if starts with "d_").
<code>show_zero_line</code>	Add horizontal line at $dZ = 0$?
<code>show_fit</code>	Add nonparametric fit?
<code>fit_method</code>	Method for fit: "loess", "gam", or "spline".

Details

In the phase diagram:

- Points where the curve crosses $dZ = 0$ are fixed points
- Negative slope at crossing indicates stability
- Positive slope indicates instability

Value

A ggplot object.

`plot_residual_diagnostics_panel`
Plot Residual Diagnostics Panel

Description

Creates a multi-panel diagnostic plot for residual analysis.

Usage

```
plot_residual_diagnostics_panel(x, ...)
```

Arguments

<code>x</code>	Object of class <code>residual_diagnostics</code>
...	Additional arguments passed to plotting functions

Value

Invisibly returns the input object

`plot_surface_3d` *3D Response Surface*

Description

Creates a 3D surface or contour plot showing how the target variable depends on two predictors.

Usage

```
plot_surface_3d(  

  data,  

  x_var,  

  y_var,  

  z_var,  

  type = c("contour", "filled_contour", "persp"),  

  n_grid = 30,  

  method = "loess"  

)
```

Arguments

data	Data frame.
x_var	First predictor variable.
y_var	Second predictor variable.
z_var	Response variable (target).
type	Plot type: "contour", "filled_contour", or "persp".
n_grid	Grid resolution for surface estimation.
method	Surface fitting method: "loess", "gam", or "linear".

Value

A plot (base graphics for persp, ggplot for contour).

plot_timeseries *Time Series Plot*

Description

Creates a time series plot with optional trend line and change point detection.

Usage

```
plot_timeseries(  
  data,  
  var,  
  time = NULL,  
  show_trend = TRUE,  
  highlight_changes = TRUE  
)
```

Arguments

data	Data frame.
var	Variable name to plot.
time	Time variable name.
show_trend	Add trend line?
highlight_changes	Highlight potential structural breaks?

Value

A ggplot object.

plot_trajectory_2d *2D Trajectory Plot*

Description

Plots the trajectory of a system in the (Z, X) plane, useful for visualizing attractors and limit cycles.

Usage

```
plot_trajectory_2d(
  data,
  x_var,
  y_var,
  time_var = NULL,
  show_arrows = TRUE,
  arrow_spacing = 10,
  show_start_end = TRUE
)
```

Arguments

<code>data</code>	Data frame.
<code>x_var</code>	First state variable.
<code>y_var</code>	Second state variable.
<code>time_var</code>	Time variable (for coloring trajectory).
<code>show_arrows</code>	Add direction arrows?
<code>arrow_spacing</code>	Spacing between arrows (every nth point).
<code>show_start_end</code>	Mark start and end points?

Value

A ggplot object.

plot_tvr_diagnostic *Diagnostic Plot for TVR Differentiation*

Description

Produces a four-panel diagnostic plot showing the original series, estimated derivative, reconstruction comparison, and residuals.

Usage

```
plot_tvr_diagnostic(Z, t = NULL, dZ_tvr)
```

Arguments

- `z` Numeric vector of original observations.
`t` Numeric vector of time points.
`dZ_tvr` TVR derivative object (from `compute_derivative_tvr`).

Value

Invisibly returns a list of diagnostic values.

`predict.variance_model`

Predict from Variance Model

Description

Predict from Variance Model

Usage

```
## S3 method for class 'variance_model'
predict(object, newdata, ...)
```

Arguments

- `object` Variance model object
`newdata` New data for prediction
`...` Additional arguments

Value

Numeric vector of predicted standard deviations.

`preprocessing`

Preprocessing Functions for Time Series Data

Description

Functions for data preparation, variable specification, and numerical differentiation including Total Variation Regularized (TVR) differentiation for noisy economic data.

print.cv_result *Print CV Results*

Description

Print CV Results

Usage

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

Arguments

x	Object of class cv_result.
...	Additional arguments (ignored).

Value

Invisibly returns the input object (called for side effects).

print.qualitative_check *Print Qualitative Check Results*

Description

Print Qualitative Check Results

Usage

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

Arguments

x	Object of class qualitative_check.
...	Additional arguments (ignored).

Value

Invisibly returns the input object (called for side effects).

```
print.residual_diagnostics
    Print Residual Diagnostics
```

Description

Print Residual Diagnostics

Usage

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

Arguments

x	Object of class residual_diagnostics
...	Additional arguments (ignored)

Value

Invisibly returns the input object (called for side effects).

```
print.tvr_derivative  Print Method for TVR Derivative
```

Description

Print Method for TVR Derivative

Usage

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

Arguments

x	A tvr_derivative object.
...	Additional arguments (ignored).

Value

Invisibly returns the input object (called for side effects).

```
print.validation_result  
      Print Validation Results
```

Description

Print Validation Results

Usage

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

Arguments

x	Object of class validation_result.
...	Additional arguments (ignored).

Value

Invisibly returns the input object (called for side effects).

```
print_summary      Print Analysis Summary
```

Description

Print Analysis Summary

Usage

```
print_summary(results)
```

Arguments

results	Analysis results list.
---------	------------------------

Value

Invisibly returns the input results object (called for side effects).

read_empirical_data *Read Empirical Data from File*

Description

Reads time series or panel data from various file formats and prepares it for use with EmpiricalDynamics functions.

Usage

```
read_empirical_data(file, time_col = NULL, date_format = NULL, ...)
```

Arguments

file	Path to the data file (CSV, RDS, or RData).
time_col	Name of the time/date column (auto-detected if NULL).
date_format	Date format string if time column is character.
...	Additional arguments passed to read.csv.

Value

A data.frame with time column converted to numeric if needed.

residual_analysis *Residual Analysis and Stochastic Differential Equations*

Description

Functions for analyzing residual structure, modeling conditional variance, constructing stochastic differential equations (SDEs), and iterative GLS estimation for heteroscedastic systems.

residual_diagnostics *Comprehensive Residual Diagnostics*

Description

Performs a battery of statistical tests on model residuals to check for autocorrelation, heteroscedasticity, and normality.

Usage

```
residual_diagnostics(
  residuals,
  data = NULL,
  predictors = NULL,
  max_lag = 10,
  plot = TRUE
)
```

Arguments

<code>residuals</code>	Numeric vector of residuals (or model object)
<code>data</code>	Optional data frame for conditional tests
<code>predictors</code>	Variable names for heteroscedasticity tests
<code>max_lag</code>	Maximum lag for autocorrelation tests
<code>plot</code>	Produce diagnostic plots?

Value

A list of test results with class "residual_diagnostics"

save_plots *Save Diagnostic Plots*

Description

Saves all diagnostic plots to files.

Usage

```
save_plots(
  results,
  output_dir,
  prefix = "empirical_dynamics",
  format = c("png", "pdf"),
  width = 8,
  height = 6,
  dpi = 300
)
```

Arguments

results	Analysis results list.
output_dir	Output directory (required, no default to comply with CRAN policy).
prefix	File name prefix.
format	Image format: "png", "pdf", "svg".
width	Plot width in inches.
height	Plot height in inches.
dpi	Resolution for raster formats.

Value

List of paths to created files.

select_equation	<i>Select Equation from Pareto Front</i>
-----------------	--

Description

Selects the best equation from the Pareto front using a specified criterion.

Usage

```
select_equation(
  results,
  criterion = c("knee", "BIC", "AIC", "min_complexity", "min_error"),
  n = NULL
)
```

Arguments

results	A symbolic_search_result object.
criterion	Selection criterion: "knee", "BIC", "AIC", "min_complexity", or "min_error".
n	Sample size (required for BIC/AIC if not stored).

Value

A symbolic_equation object.

`select_lambda_cv_tvr` *Cross-Validation Selection of Lambda for TVR*

Description

Selects the regularization parameter lambda using leave-one-out-like cross-validation.

Usage

```
select_lambda_cv_tvr(
  Z,
  t,
  A = NULL,
  D = NULL,
  solver = "osqp",
  lambda_seq = 10^seq(-4, 2, length.out = 30),
  verbose = FALSE
)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.
A	Integration matrix.
D	Difference matrix.
solver	Optimization backend.
lambda_seq	Sequence of lambda values to evaluate.
verbose	Print progress?

Value

Selected lambda value.

sensitivity_analysis *Parameter Sensitivity Analysis*

Description

Examines how sensitive the model predictions are to parameter perturbations.

Usage

```
sensitivity_analysis(  
  equation,  
  data,  
  response = NULL,  
  derivative_col = NULL,  
  perturbation_pct = 10,  
  n_bootstrap = 100  
)
```

Arguments

equation	Fitted equation object.
data	Data for evaluation.
response	Name of response column.
derivative_col	Alias for response.
perturbation_pct	Percentage perturbation (default 10%).
n_bootstrap	Number of bootstrap samples for uncertainty.

Value

Data frame with sensitivity metrics for each parameter.

setup_julia_backend *Setup Julia Backend*

Description

Checks if Julia and the required SymbolicRegression.jl package are installed.

Usage

```
setup_julia_backend()
```

Value

Logical indicating if the backend is ready.

`simulate_trajectory` *Simulate Trajectory from SDE*

Description

Simulates trajectories using the discovered SDE to assess whether the model can reproduce observed dynamics.

Usage

```
simulate_trajectory(
  sde,
  initial_conditions,
  times,
  n_sims = 100,
  method = c("euler", "milstein", "rk4"),
  exogenous_data = NULL,
  seed = NULL
)
```

Arguments

<code>sde</code>	SDE object from <code>construct_sde</code> or <code>estimate_sde_iterative</code> .
<code>initial_conditions</code>	Named vector of initial values for all variables.
<code>times</code>	Numeric vector of time points.
<code>n_sims</code>	Number of Monte Carlo simulations (for stochastic models).
<code>method</code>	Integration method: "euler", "milstein", "rk4" (deterministic only).
<code>exogenous_data</code>	Data frame with exogenous variable trajectories (if any).
<code>seed</code>	Random seed for reproducibility.

Value

Object of class "trajectory_simulation" containing:

<code>trajectories</code>	Array of simulated trajectories (time x variable x simulation)
<code>times</code>	Time points
<code>summary</code>	Summary statistics (mean, quantiles) at each time

Examples

```
# Toy example: dX = 0.5 * X
# Mock SDE object structure
sde <- list(
  drift = list(expression = "0.5 * X"),
  diffusion = list(expression = "0.1"), # Add noise
```

```

    variable = "X"
)
class(sde) <- "sde_model"

# Simulation
sim <- simulate_trajectory(
  sde = sde,
  initial_conditions = c(X = 1),
  times = seq(0, 1, by = 0.1),
  n_sims = 10,
  seed = 123
)
print(sim$summary$mean)

```

specify_variables *Specify Variable Types for Dynamical Analysis*

Description

Classifies variables in a dataset according to their role in the dynamical system being studied.

Usage

```

specify_variables(
  data,
  endogenous = NULL,
  endogenous_coupled = NULL,
  coupled = NULL,
  exogenous = NULL,
  slow_parameter = NULL,
  time = NULL,
  time_col = NULL
)

```

Arguments

<code>data</code>	A <code>data.frame</code> containing the time series data.
<code>endogenous</code>	Character vector of endogenous state variable names.
<code>endogenous_coupled</code>	Character vector of coupled endogenous variables.
<code>coupled</code>	Alias for <code>endogenous_coupled</code> (for compatibility).
<code>exogenous</code>	Character vector of exogenous forcing variable names.
<code>slow_parameter</code>	Character vector of slowly-varying parameter names.
<code>time</code>	Name of the time column (auto-detected if <code>NULL</code>).
<code>time_col</code>	Alias for <code>time</code> (for compatibility).

Details

Variable types:

- **endogenous**: Variables whose dynamics are modeled (appear as dZ/dt).
- **endogenous_coupled**: Variables that co-evolve with endogenous vars.
- **exogenous**: Variables that influence the system but are not modeled.
- **slow_parameter**: Variables that change on much longer timescales.

Value

The input data.frame with variable specifications added as the "var_spec" attribute.

Examples

```
data <- data.frame(
  time = 1:10,
  profit_rate = runif(10),
  capital_stock = runif(10),
  interest_rate = runif(10)
)

data <- specify_variables(data,
  endogenous = "profit_rate",
  endogenous_coupled = "capital_stock",
  exogenous = "interest_rate"
)
attr(data, "var_spec")
```

suggest_differentiation_method

Suggest Differentiation Method Based on Data Characteristics

Description

Analyzes the time series to recommend the most appropriate differentiation method based on detected features like trend, periodicity, shocks, and noise.

Usage

```
suggest_differentiation_method(Z, t = NULL)
```

Arguments

Z	Numeric vector of observations.
t	Numeric vector of time points.

Value

List with suggested method and diagnostic information.

Examples

```
t <- 1:100
Z <- 0.1 * t + rnorm(100) # Trend with noise
result <- suggest_differentiation_method(Z, t)
print(result$suggested_method)
```

Description

Functions for discovering functional forms through symbolic regression using genetic algorithms. Interfaces with Julia's `SymbolicRegression.jl` for advanced search, with fallback to R-native methods for simpler cases.

Discovers the functional form of a differential equation from data using genetic/evolutionary algorithms. Returns a Pareto front of equations trading off complexity against fit.

Usage

```
symbolic_search(
  target,
  predictors,
  operators = NULL,
  constraints = NULL,
  n_runs = 5,
  complexity_penalty = 0.05,
  parsimony_pressure = c("adaptive", "constant", "none"),
  backend = c("r_genetic", "julia", "r_exhaustive"),
  julia_options = NULL,
  weights = NULL,
  verbose = TRUE
)
```

Arguments

<code>target</code>	Numeric vector of target values (typically derivatives).
<code>predictors</code>	Data frame of predictor variables.
<code>operators</code>	List specifying allowed operators: <ul style="list-style-type: none"> • binary: <code>c("+", "-", "*", "/")</code> • unary: <code>c("exp", "log", "sqrt", "inv", "square")</code> • custom: Custom function names (must be defined)

<code>constraints</code>	List of constraints:
	<ul style="list-style-type: none"> • forced: Formula of terms that must appear • forbidden: Formula of terms that must not appear • max_complexity: Maximum expression complexity
<code>n_runs</code>	Number of independent runs for robustness.
<code>complexity_penalty</code>	Penalty per unit complexity.
<code>parsimony_pressure</code>	Type of parsimony: "constant", "adaptive", or "none".
<code>backend</code>	Computation backend: "julia", "r_genetic", or "r_exhaustive".
<code>julia_options</code>	List of options passed to SymbolicRegression.jl.
<code>weights</code>	Optional weight vector for weighted regression.
<code>verbose</code>	Print progress messages?

Value

An object of class "symbolic_search_result" containing:

- `pareto_front`: Data frame of Pareto-optimal equations
- `all_equations`: All discovered equations
- `best_by_complexity`: Best equation at each complexity level
- `run_diagnostics`: Information about each run

Examples

```
# Toy example using R-native exhaustive search (fastest for demo)
data <- data.frame(
  x = seq(1, 10, length.out = 20),
  y = seq(1, 10, length.out = 20)^2 + rnorm(20, sd = 0.1)
)

# Discover y ~ x^2
results <- symbolic_search(
  target = data$y,
  predictors = data["x"],
  backend = "r_exhaustive"
)

print(head(results$pareto_front))
```

symbolic_search_weighted
Weighted Symbolic Search

Description

Performs symbolic search with weighted least squares.

Usage

```
symbolic_search_weighted(  
  target,  
  predictors,  
  weights,  
  operators = NULL,  
  constraints = NULL,  
  n_runs = 3,  
  complexity_penalty = 0.05,  
  verbose = TRUE  
)
```

Arguments

target	Numeric vector of target values (typically derivatives).
predictors	Data frame of predictor variables.
weights	Optional weight vector for weighted regression.
operators	List specifying allowed operators: <ul style="list-style-type: none">• binary: c("+", "-", "*", "/")• unary: c("exp", "log", "sqrt", "inv", "square")• custom: Custom function names (must be defined)
constraints	List of constraints: <ul style="list-style-type: none">• forced: Formula of terms that must appear• forbidden: Formula of terms that must not appear• max_complexity: Maximum expression complexity
n_runs	Number of independent runs for robustness.
complexity_penalty	Penalty per unit complexity.
verbose	Print progress messages?

Value

A symbolic_search_result object

to_latex*Convert Equation to LaTeX***Description**

Converts a discovered equation to LaTeX format for publication.

Usage

```
to_latex(
  equation,
  variable = "Z",
  precision = 3,
  scientific_notation = TRUE,
  include_uncertainty = FALSE,
  se_values = NULL
)
```

Arguments

<code>equation</code>	Fitted equation object.
<code>variable</code>	Name of the dependent variable (for dZ/dt notation).
<code>precision</code>	Number of decimal places for coefficients.
<code>scientific_notation</code>	Use scientific notation for large/small coefficients.
<code>include_uncertainty</code>	Include standard errors in parentheses.
<code>se_values</code>	Named vector of standard errors (optional).

Value

Character string with LaTeX equation.

Examples

```
# Toy example using a linear model
data <- data.frame(Z = 1:10, dZ = 2 * (1:10) + 3)
model <- stats::lm(dZ ~ Z, data = data)

# Convert to LaTeX
latex_eq <- to_latex(model, variable = "Z")
cat(latex_eq)
```

Description

Runs a battery of validation tests on the discovered equation.

Usage

```
validate_model(  
  equation,  
  sde = NULL,  
  data,  
  response = NULL,  
  derivative_col = NULL,  
  variable,  
  time_col = "time",  
  cv_folds = 5,  
  n_sims = 50,  
  expected_features = list(),  
  verbose = TRUE  
)
```

Arguments

equation	Fitted equation object.
sde	SDE object (optional, for trajectory validation).
data	Original data frame.
response	Name of response column.
derivative_col	Alias for response (for compatibility).
variable	Main variable name.
time_col	Time column name.
cv_folds	Number of CV folds.
n_sims	Number of trajectory simulations.
expected_features	List of expected qualitative features.
verbose	Print progress.

Value

Object of class "validation_result".

validation*Validation of Discovered Equations*

Description

Functions for validating discovered differential equations through cross-validation, trajectory simulation, and qualitative behavior analysis.

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