

# Package ‘EmpiricalDynamics’

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**Title** Empirical Discovery of Differential Equations from Time Series  
Data

**Version** 0.1.3

**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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**NeedsCompilation** no

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---

analysis_summary	<i>Create Analysis Summary</i>
------------------	--------------------------------

---

**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

<code>equation</code>	Fitted equation object.
<code>variable</code>	Name of the main variable.
<code>range</code>	Numeric vector of length 2 specifying search range.
<code>n_grid</code>	Number of grid points for initial search.
<code>exogenous_values</code>	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

data                    Data frame (with exploration results as attribute).  
hypotheses            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: <ul style="list-style-type: none"> <li>• n_fixed_points: Expected number of fixed points</li> <li>• stability_pattern: e.g., c("stable", "unstable", "stable")</li> <li>• monotonicity: Expected sign of derivative ("positive", "negative", "none")</li> <li>• bounded: Whether dynamics should be bounded</li> </ul>

**Value**

Object of class "qualitative\_check".

---

coefficient_table	<i>Generate Coefficient Table</i>
-------------------	-----------------------------------

---

**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**

simulation	Trajectory simulation object.
observed_data	Data frame with observed values.
time_col	Name of time column.
var_col	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

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

### 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

Z	Numeric vector of observations.
t	Numeric vector of time points.
...	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. The problem is internally rescaled for numerical stability, and a cascading solver chain is used for maximum robustness.

**Usage**

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

**Arguments**

Z	Numeric vector of observations.
t	Numeric vector of time points (NULL assumes dt=1).
lambda	Regularization parameter ("auto" for cross-validation selection).
solver	Preferred solver: "clarabel" (default, interior point), "scs" (conic ADMM), or "osqp" (QP ADMM). If the preferred solver fails or returns an inaccurate solution, other solvers are tried automatically.
...	Additional arguments (ignored).

**Details****Solver status interpretation:**

- "optimal": The solver converged within its tolerances. Results are reliable.
- "optimal\_inaccurate": The solver found a solution but did not meet strict tolerance criteria. For TVR this rarely affects derivative quality in a perceptible way, because internal rescaling keeps the problem well-conditioned. If degradation is observed, try a different preferred solver or adjust lambda manually.

**TVR and stochastic differential equations:** TVR penalizes total variation of the derivative, which effectively smooths out high-frequency components. In SDEs, the diffusion signature lives in those high-frequency residuals. As a consequence, the better TVR estimates the drift, the more it removes the information needed to recover the diffusion coefficient from residuals. For SDE applications, consider using [estimate\\_diffusion\\_qv](#) (quadratic variation) to estimate diffusion directly from increments rather than from TVR residuals.

**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**

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

**Value**

Numeric vector of residuals

---

construct_sde	<i>Construct Stochastic Differential Equation Model</i>
---------------	---

---

**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**

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

<code>data</code>	Data frame.
<code>transformations</code>	List of formulas specifying transformations.
<code>variables</code>	Character vector of variable names (alternative interface).
<code>cols</code>	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      *Cross-Validate Discovered Equation*

---

### 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

equation	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> .
data	Data frame containing all variables.
response	Name of the response column (derivative).
derivative_col	Alias for response (for compatibility).
k	Number of folds for cross-validation.
method	CV method: "random", "block", "rolling".
block_size	For block methods, size of contiguous blocks.
horizon	For rolling CV, forecast horizon.
refit_derivative	Logical; whether to recompute derivatives for each fold (currently unused).
diff_method	Differentiation method if refitting (currently unused).
verbose	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\_diffusion\_qv *Estimate Diffusion via Quadratic Variation*

---

**Description**

Estimates the diffusion coefficient  $g(\cdot)$  of an SDE directly from the quadratic variation of increments, without relying on derivative residuals. For an SDE  $dZ = f dt + g dW$ , the realized quadratic variation satisfies  $E[(\Delta Z)^2/\Delta t] = g^2 + O(\Delta t)$ , providing a model-free estimator of  $g^2$ .

**Usage**

```
estimate_diffusion_qv(
  Z,
  t,
  predictors,
  data = NULL,
  method = c("linear", "quadratic", "gam", "constant"),
  smooth_window = 21L
)
```

**Arguments**

Z	Numeric vector of the state variable.
t	Numeric vector of time points.
predictors	Character vector of predictor names or data frame.
data	Data frame containing the predictors (required if predictors is a character vector).
method	Modeling method for $g^2$ : "linear", "quadratic", "gam", or "constant".
smooth_window	Size of the rolling median window for smoothing raw quadratic variation (must be odd, default 21).

**Details**

This method is recommended over residual-based diffusion estimation when TVR is used to compute derivatives. TVR smooths out high-frequency components that carry the diffusion signature, making residual-based estimation unreliable. Quadratic variation bypasses this by working directly with the raw increments.

The raw per-step estimates  $(\Delta Z)^2/\Delta t$  are noisy (each is a single chi-squared realization), so a rolling median is applied before fitting.

**Value**

An object of class "variance\_model" compatible with the SDE pipeline.

---

`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**

<code>expression</code>	Character string with the equation expression.
<code>data</code>	Data frame with variables.
<code>response</code>	Name of the response variable.
<code>derivative_col</code>	Alias for response (for compatibility).
<code>method</code>	Estimation method: "grid_search", "random", or "heuristic".
<code>n_tries</code>	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,
  diffusion_method = c("quadratic_variation", "residual"),
  Z = NULL,
  t = NULL
)
```

### Arguments

target	Numeric vector of target values (derivatives)
predictors	Data frame of predictor variables
data	Full data frame
initial_drift	Initial drift equation (optional)
max_iter	Maximum number of iterations
tol	Convergence tolerance (RMSE change in coefficients)
diffusion_method	Method for estimating the final diffusion coefficient: "quadratic_variation" (default) estimates $g(\cdot)$ directly from increments $(\Delta Z)^2/dt$ , which is robust when TVR is used for derivatives. "residual" uses the classical residual-based approach via <a href="#">model_conditional_variance</a> .
Z	Numeric vector of the state variable (required when diffusion_method = "quadratic_variation").
t	Numeric vector of time points (required when diffusion_method = "quadratic_variation").

### Value

An sde\_model object with refined estimates

---

exploration	<i>Visual Exploration of Dynamical Structure</i>
-------------	--

---

### 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.

---

explore_dynamics	<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

data	Data frame containing the time series.
target	Name of the target variable (or its derivative).
predictors	Character vector of predictor variable names.
time	Name of the time column (auto-detected if NULL).
n_bins	Number of bins for conditional analysis.
include	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**

results	Analysis results list.
output_dir	Output directory (required, no default to comply with CRAN policy).
prefix	File name prefix.
formats	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

*Format Equation for Display*

---

## 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

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

---

**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	<i>List Available Example Datasets</i>
-------------------	--

---

**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

name                      Name of the dataset to load. Available datasets:

- "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**

<code>equations</code>	Named list of fitted equation objects.
<code>data</code>	Data for computing fit statistics.
<code>derivative_col</code>	Response variable column.
<code>format</code>	Output format.
<code>caption</code>	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**

residuals	Numeric vector of residuals
predictors	Formula or data frame of predictor variables (or vector of names)
data	Data frame (if predictors is a formula or vector of names)
method	Modeling method: "symbolic", "linear", "quadratic", "gam", or "constant"
transform	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	<i>Output and Report Generation</i>
--------	-------------------------------------

---

**Description**

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

---

plot.bifurcation_analysis	<i>Plot Bifurcation Diagram</i>
---------------------------	---------------------------------

---

**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	<i>Plot CV Results</i>
----------------	------------------------

---

**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	<i>Plot Simulated Trajectories</i>
----------------------------	------------------------------------

---

**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**

data	Data frame.
x_var	X variable name.
y_var	Y variable name.
color_var	Optional variable for color mapping.
show_fit	Add smooth fit?
show_marginals	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**

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

**Value**

A ggplot object.

---

plot_phase_1d	<i>1D Phase Diagram</i>
---------------	-------------------------

---

### 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>
<code>...</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	<i>Time Series Plot</i>
-----------------	-------------------------

---

**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

data	Data frame.
x_var	First state variable.
y_var	Second state variable.
time_var	Time variable (for coloring trajectory).
show_arrows	Add direction arrows?
arrow_spacing	Spacing between arrows (every nth point).
show_start_end	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

residuals	Numeric vector of residuals (or model object)
data	Optional data frame for conditional tests
predictors	Variable names for heteroscedasticity tests
max_lag	Maximum lag for autocorrelation tests
plot	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. Problems are internally rescaled for numerical stability, and a solver chain is used for each candidate lambda.

**Usage**

```
select_lambda_cv_tvr(
  Z,
  t,
  A = NULL,
  D = NULL,
  solver = "clarabel",
  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	Preferred solver (default "clarabel"). Falls back to other solvers if the preferred one fails.
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

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

### Value

Object of class "trajectory\_simulation" containing:

trajectories	Array of simulated trajectories (time x variable x simulation)
times	Time points
summary	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

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

---

symbolic\_search

*Symbolic Regression and Equation Discovery*


---

**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**

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

constraints	List of constraints: <ul style="list-style-type: none"> <li>• forced: Formula of terms that must appear</li> <li>• forbidden: Formula of terms that must not appear</li> <li>• max_complexity: Maximum expression complexity</li> </ul>
n_runs	Number of independent runs for robustness.
complexity_penalty	Penalty per unit complexity.
parsimony_pressure	Type of parsimony: "constant", "adaptive", or "none".
backend	Computation backend: "julia", "r_genetic", or "r_exhaustive".
julia_options	List of options passed to SymbolicRegression.jl.
weights	Optional weight vector for weighted regression.
verbose	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  $y \sim x^2$ 
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"><li>• binary: c("+", "-", "*", "/")</li><li>• unary: c("exp", "log", "sqrt", "inv", "square")</li><li>• custom: Custom function names (must be defined)</li></ul>
constraints	List of constraints: <ul style="list-style-type: none"><li>• forced: Formula of terms that must appear</li><li>• forbidden: Formula of terms that must not appear</li><li>• max_complexity: Maximum expression complexity</li></ul>
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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