

# Package ‘MultiSpline’

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**Type** Package

**Title** Spline-Based Nonlinear Modeling for Multilevel and Longitudinal Data

**Version** 0.1.0

**Description** Provides tools for fitting, predicting, and visualizing nonlinear relationships in single-level, multilevel, and longitudinal regression models. Flexible functional forms are supported using natural cubic splines via 'splines' and smooth terms via 'mgcv'. The package offers a unified interface for specifying nonlinear effects, interactions with time variables, clustering through random intercepts, and additional linear covariates. Utilities are included to generate prediction grids and produce effect plots, facilitating interpretation and visualization of nonlinear effects in applied regression workflows.

**Depends** R (>= 4.2.0)

**Imports** stats, lme4, mgcv, dplyr, ggplot2, rlang

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**nl\_fit***Fit a nonlinear (spline or GAM) single-level or multilevel model***Description**

Fits a nonlinear regression model for an outcome *y* with a focal predictor *x*, modeled either by a natural cubic spline (`splines::ns()`) or a GAM smooth (`mgcv::s()`).

Optionally includes a time variable *time*. For spline fits (`method = "ns"`), multilevel random-intercept structure can be added via one or more grouping variables in `cluster`.

Models fitted:

- Single-level spline: `stats::lm()`
- Single-level GAM: `mgcv::gam()`
- Multilevel spline: `lme4::lmer()` (Gaussian) or `lme4::glmer()` (non-Gaussian)

**Usage**

```
nl_fit(
  data,
  y,
  x,
  time = NULL,
  cluster = NULL,
  controls = NULL,
  method = c("ns", "gam"),
  df = 4,
  k = 5,
  family = stats::gaussian(),
  ...
)
```

**Arguments**

<code>data</code>	A data frame (often long format for longitudinal data).
<code>y</code>	Outcome variable name (string).
<code>x</code>	Focal nonlinear predictor name (string). Must be numeric.
<code>time</code>	Optional time variable name (string). If provided, a spline-by-time interaction is included for <code>method = "ns"</code> ; for <code>method = "gam"</code> , a factor <code>time</code> uses <code>s(x, by = time)</code> (group-specific smooths), while a numeric <code>time</code> uses <code>s(x, k = k) + time + ti(x, time, k = k)</code> (tensor interaction).
<code>cluster</code>	Optional character vector of grouping variable name(s) for random intercepts, e.g., <code>NULL</code> , <code>"id"</code> , or <code>c("id", "schid")</code> .
<code>controls</code>	Optional character vector of additional covariate names to include linearly.
<code>method</code>	Either <code>"ns"</code> (natural spline) or <code>"gam"</code> (GAM smooth). Multilevel fits are currently supported only for <code>"ns"</code> .
<code>df</code>	Degrees of freedom for <code>splines::ns()</code> when <code>method = "ns"</code> . Must be a single numeric value $\geq 1$ . Default is 4.

<code>k</code>	Basis dimension for <code>mgcv::s()</code> and <code>mgcv::ti()</code> when <code>method = "gam"</code> . Must be a single numeric value $\geq 3$ . Default is 5.
<code>family</code>	A model family object, e.g., <code>stats::gaussian()</code> or <code>stats::binomial()</code> . For multilevel fits, <code>gaussian()</code> uses <code>lme4::lmer()</code> and non-Gaussian families use <code>lme4::glmer()</code> .
<code>...</code>	Additional arguments passed to the underlying fitting function ( <code>stats::lm()</code> , <code>mgcv::gam()</code> , <code>lme4::lmer()</code> , or <code>lme4::glmer()</code> ).

### Value

An object of class "`nl_fit`" (a named list) containing:

- `model` The fitted model object.
- `method` The fitting method used ("ns" or "gam").
- `y` Name of the outcome variable.
- `x` Name of the focal predictor.
- `time` Name of the time variable, or NULL.
- `cluster` Character vector of clustering variables, or NULL.
- `controls` Character vector of control variable names, or NULL.
- `df` Degrees of freedom used for `splines::ns()`.
- `k` Basis dimension used for `mgcv::s()` / `mgcv::ti()`.
- `family` The family object used.
- `formula` The model formula passed to the fitter.
- `call` The matched call.
- `x_info` A list with quantiles and range of `x` for building prediction grids: `q01`, `q99`, `min`, `max`, `n`.
- `levels_info` A named list of factor levels for `time` and any factor control variables.
- `control_defaults` A named list of typical values for control variables: the mean for numeric variables and the first level for factors.

### See Also

[ns](#), [gam](#), [lmer](#), [glmer](#)

### Examples

```
## Not run:
# Single-level natural spline
fit_sl <- nl_fit(data = mydata, y = "outcome", x = "age", df = 4)

# Single-level GAM
fit_gam <- nl_fit(
  data    = mydata,
  y       = "outcome",
  x       = "age",
  method  = "gam",
  k       = 5
)

# Multilevel spline with random intercepts
fit_ml <- nl_fit(
```

```

data      = mydata,
y        = "outcome",
x        = "age",
cluster  = "id",
df       = 4
)

# Spline with time interaction and controls
fit_t <- nl_fit(
  data      = mydata,
  y        = "outcome",
  x        = "age",
  time    = "wave",
  controls = c("sex", "baseline_score"),
  df       = 4
)

## End(Not run)

```

**nl\_icc***Intraclass correlation coefficients for a multilevel nl\_fit model***Description**

Extracts variance components from a multilevel `nl_fit` object and computes intraclass correlation coefficients (ICCs) for each grouping level plus the residual.

The ICC for grouping factor  $g$  is defined as:

$$ICC_g = \frac{\sigma_g^2}{\sum_j \sigma_j^2 + \sigma_\epsilon^2}$$

**Usage**

```
nl_icc(object, include_residual = TRUE)
```

**Arguments**

<code>object</code>	An <code>nl_fit</code> object returned by <code>nl_fit</code> that was fitted with one or more cluster variables (i.e., a multilevel model fitted via <code>lme4::lmer()</code> or <code>lme4::glmer()</code> ).
<code>include_residual</code>	Logical; if TRUE (default), includes the residual variance component <code>ICC_resid</code> in the output so that all values sum to 1.

**Value**

A named numeric vector of ICCs, one per grouping factor (named `ICC_<groupname>`) plus `Residual` for the residual variance (when `include_residual = TRUE`). All values sum to 1.

**See Also**

`nl\_fit`

## Examples

```
## Not run:
fit <- nl_fit(
  data      = mydata,
  y         = "math_score",
  x         = "SES",
  cluster   = c("id", "schid"),
  df        = 4
)
nl_icc(fit)

## End(Not run)
```

nl\_plot

*Plot predictions from nl\_predict()*

## Description

Visualizes predicted nonlinear effects returned by [nl\\_predict](#). If a time variable is present (or detected), draws separate colored curves for each time level with optional confidence ribbons.

## Usage

```
nl_plot(
  pred_df,
  x,
  time = NULL,
  show_ci = TRUE,
  ci_level = 0.95,
  y_lab = "Predicted outcome",
  x_lab = NULL,
  title = NULL,
  legend_title = NULL
)
```

## Arguments

<code>pred_df</code>	A data frame returned by <a href="#">nl_predict</a> . Must contain at minimum columns fit and the focal predictor <code>x</code> . For confidence bands, provide either <code>lwr/upr</code> or <code>se.fit</code> .
<code>x</code>	Character string naming the focal predictor column in <code>pred_df</code> .
<code>time</code>	Optional character string naming the time variable column in <code>pred_df</code> . If <code>NULL</code> , the function will try to auto-detect a sensible time column (e.g., "TimePoint", "time", "wave") if present.
<code>show_ci</code>	Logical; if <code>TRUE</code> , adds a confidence ribbon. Default <code>TRUE</code> .
<code>ci_level</code>	Numeric in $(0, 1)$ ; confidence level used when computing CI from <code>se.fit</code> . Default $0.95$ .
<code>y_lab</code>	Character label for the y-axis. Default "Predicted outcome".
<code>x_lab</code>	Character label for the x-axis. Defaults to the value of <code>x</code> .
<code>title</code>	Optional plot title string.
<code>legend_title</code>	Optional legend title string. Defaults to the value of <code>time</code> .

**Value**

A ggplot object.

**See Also**

[nl\\_predict](#), [nl\\_fit](#)

[nl\\_predict](#)

*Generate predictions from an nl\_fit model*

**Description**

Creates a prediction data frame over a grid of the focal predictor `x` (and optionally over `time`), holding control variables at typical values (means for numeric; reference levels for factors). For mixed models, predictions default to population-level curves (random effects excluded).

**Usage**

```
nl_predict(
  object,
  x_seq = NULL,
  time_levels = NULL,
  controls_fixed = NULL,
  se = TRUE,
  level = 0.95,
  re_form = NA,
  ...
)
```

**Arguments**

<code>object</code>	An <code>nl_fit</code> object returned by <a href="#">nl_fit</a> .
<code>x_seq</code>	Optional numeric vector of <code>x</code> values to predict over. If <code>NULL</code> , uses 100 evenly-spaced points between the stored 1st and 99th percentiles (with fallback to the stored min/max range).
<code>time_levels</code>	Optional vector of time levels to predict for. If <code>NULL</code> and a time variable is present, uses stored factor levels.
<code>controls_fixed</code>	Optional named list giving specific values for control variables. If <code>NULL</code> , stored defaults are used (mean for numeric; first level for factors).
<code>se</code>	Logical; if <code>TRUE</code> , includes standard errors and confidence intervals where available. Default <code>TRUE</code> .
<code>level</code>	Confidence level for the interval. Default <code>0.95</code> .
<code>re_form</code>	For mixed models ( <code>lmerMod</code> / <code>glmerMod</code> ), passed to <code>predict()</code> . Default <code>NA</code> gives population-level predictions (random effects set to zero).
<code>...</code>	Reserved for future use.

**Value**

A data frame (or tibble) with columns: the focal predictor `x`, `time` (if applicable), any control variables (at fixed values), `fit`, `se.fit`, `lwr`, and `upr`.

**See Also**[nl\\_fit](#), [nl\\_plot](#)

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**nl\_summary***Tidy coefficient table for an nl\_fit model*

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**Description**

Produces a tidy coefficient table for a model fitted by [nl\\_fit](#). For linear mixed models (lmerMod), optional p-values and denominator degrees of freedom are obtained via **ImerTest** (Satterthwaite method). For GLMMs (glmerMod), z-tests are reported from `summary()`.

**Usage**

```
nl_summary(  
  object,  
  digits = 3,  
  pvals = TRUE,  
  df_method = c("satterthwaite", "none")  
)
```

**Arguments**

<code>object</code>	An <code>nl_fit</code> object returned by <a href="#">nl_fit</a> .
<code>digits</code>	Integer; number of decimal places for rounding. If NULL, no rounding is applied. Default 3.
<code>pvals</code>	Logical; if TRUE, attempts to include p-values. Default TRUE.
<code>df_method</code>	For lmerMod with <code>pvals = TRUE</code> : "satterthwaite" (requires <b>ImerTest</b> ) or "none". Default "satterthwaite".

**Value**

A data frame (or tibble) with columns: Term, Estimate, Std.Error, df (if available), statistic, and p.value (if available).

**See Also**[nl\\_fit](#), [summary.nl\\_fit](#)

**print.nl\_fit** *Print method for nl\_fit objects*

## Description

Compact console display for objects returned by [nl\\_fit](#). Shows key metadata (method, outcome, predictor, time, clustering, family, and controls) and the fitted model formula.

## Usage

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

## Arguments

- x An object of class `nl_fit`.
- ... Further arguments (currently ignored).

## Value

x invisibly.

**print.summary\_nl\_fit** *Print method for summary\_nl\_fit objects*

## Description

Prints a `summary_nl_fit` object returned by [summary.nl\\_fit](#).

## Usage

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

## Arguments

- x A `summary_nl_fit` object.
- ... Further arguments passed to `print`.

## Value

x invisibly.

---

summary.nl_fit	<i>Summary method for nl_fit objects</i>
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## Description

Produces a tidy coefficient table via [nl\\_summary](#) and wraps it in a `summary_nl_fit` object for pretty printing.

## Usage

```
## S3 method for class 'nl_fit'  
summary(  
  object,  
  digits = 3,  
  pvals = TRUE,  
  df_method = c("satterthwaite", "none"),  
  ...  
)
```

## Arguments

object	An <code>nl_fit</code> object returned by <a href="#">nl_fit</a> .
digits	Number of decimal places for rounding. Default 3.
pvals	Logical; if TRUE, attempts to include p-values. Default TRUE.
df_method	For <code>lmerMod</code> : "satterthwaite" (requires <a href="#">lmerTest</a> ) or "none". Default "satterthwaite".
...	Further arguments passed to <a href="#">nl_summary</a> .

## Value

An object of class `summary_nl_fit` containing `call`, `formula`, `method`, and `table`.

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