

Package: `poissonsUPERlearner` (via `r-universe`)

May 18, 2026

Title Poisson Super Learner

Version 0.2.0

Description Provides tools for fitting piecewise-constant hazard models for survival and competing risks data, including ensemble hazard estimation via the Super Learner framework. The package supports estimation of survival functions and absolute risk predictions from fitted cause-specific hazard models. For the Super Learner framework see van der Laan, Polley and Hubbard (2007) <[doi:10.2202/1544-6115.1309](https://doi.org/10.2202/1544-6115.1309)>.

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LinkingTo Rcpp

Encoding UTF-8

RoxygenNote 7.3.3

Depends data.table, sampling, riskRegression

Imports Rcpp, methods, lava, Matrix, glmnet, mgcv

Suggests knitr, rmarkdown, survival, prodlim, testthat (>= 3.0.0)

VignetteBuilder knitr

Config/testthat/edition 3

NeedsCompilation yes

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Repository <https://cran.r-universe.dev>

Date/Publication 2026-05-18 19:09:28 UTC

RemoteUrl <https://github.com/cran/poissonsUPERlearner>

RemoteRef HEAD

RemoteSha 032b486e2cf2cd63cb1733c0174bfe6ad8c65c6c

Contents

coef.base_learner	2
coef.poisson_superlearner	3
fit_learner	5
Learner_gam-class	7
Learner_glmnet-class	8
Learner_hal-class	9
pch_absolute_risk	11
pch_absolute_risk_euler	12
pch_survival	13
predict.base_learner	14
predict.poisson_superlearner	16
predictRisk.base_learner	18
predictRisk.poisson_superlearner	19
print.base_learner	20
print.poisson_superlearner	21
simulateStenoT1	22
summary.base_learner	25
summary.poisson_superlearner	26
Superlearner	27
Index	31

coef.base_learner	<i>Extract coefficients from a fitted base learner</i>
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Description

Convenience method to extract (cause-specific) model coefficients from a fitted `base_learner` returned by `fit_learner()`.

Usage

```
## S3 method for class 'base_learner'
coef(object, cause = NULL, ...)
```

Arguments

<code>object</code>	<code>base_learner</code> . A fitted object returned by <code>fit_learner()</code> .
<code>cause</code>	<code>numeric(1)</code> or <code>NULL</code> . Which cause to extract coefficients for. If <code>NULL</code> , coefficients are returned for all causes. Causes are indexed 1, 2, ..., <code>object\$data_info\$n_crisks</code> (with 0 reserved for censoring).
<code>...</code>	Passed to the underlying <code>coef()</code> method of the fitted learner object (learner-dependent; e.g., <code>s</code> for <code>glmnet</code>).

Details

For competing risks, `fit_learner()` fits one model per cause, stored in `object$learner_fit[[k]]` for $k = 1, 2, \dots, K$. This method simply dispatches to the underlying model's `coef()` method for each fitted object.

Learner-dependent output. The returned coefficient object depends on the base learner used (e.g. a numeric vector, a sparse matrix, a list, etc.). This method does not post-process or rename coefficients; it returns the output of `coef(object$learner_fit[[k]], ...)` unchanged.

Value

If `cause` is a single integer, returns the coefficient object produced by `coef()` for that cause-specific fitted model.

If `cause = NULL`, returns a list of length `object$data_info$n_crisks`, where element `[[k]]` contains coefficients for cause k .

If no fitted model is present (`object$learner_fit` is `NULL`), signals a message and returns `invisible(object)`.

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)
lrn <- Learner_glmnet(covariates = c("age", "value_LDL"),
                    lambda = 0, cross_validation = FALSE)
bl <- fit_learner(d, learner = lrn, id = "id",
                status = "status_cvd", event_time = "time_cvd",
                number_of_nodes = 4)

# coefficients for cause 1
coef(bl, cause = 1)

# coefficients for all causes (list)
coef(bl)
```

```
coef.poisson_superlearner
```

Extract stacking (meta-learner) coefficients from a fitted Poisson Super Learner

Description

Extracts the **meta-learner coefficients** (stacking weights) from a fitted `poisson_superlearner` object returned by `Superlearner()`.

Usage

```
## S3 method for class 'poisson_superlearner'
coef(object, cause = NULL, model = "sl", ...)
```

Arguments

object	poisson_superlearner. A fitted ensemble returned by <code>Superlearner()</code> .
cause	numeric(1) or NULL. Which cause to extract meta-learner coefficients for. If NULL, coefficients are returned for all causes. Causes are indexed 1, 2, ..., <code>object\$data_info\$n_crisis</code> .
model	Model selector. Default is "s1" for the stacked super learner. Allowed values are: <ul style="list-style-type: none"> 0, "s1", "superlearner", or "super_learner" Extract coefficients from the stacked meta-learner. For causes with no fitted meta-learner, this falls back to the retained base learner. "discrete_s1" and aliases Extract coefficients from the cause-specific base learners with the smallest cross-validated deviance. learner label Extract coefficients from one stored base learner by its label in <code>object\$data_info\$learners_labels[[k]]</code>. "learner_j" or character integer "j" Extract coefficients from the j-th stored learner. integer j $j \geq 1$ Extract coefficients from the j-th stored learner. vector of labels or positive integer indices Use cause-specific base learners; length must equal <code>object\$data_info\$n_crisis</code>.
...	Passed to the underlying <code>coef()</code> method of the fitted meta-learner (learner-dependent; e.g., <code>s</code> for <code>glmnet</code>).

Details

For each cause k , the ensemble stores a fitted meta-learner in `object$superlearner[[k]]$meta_learner_fit`. This method dispatches to the underlying `coef()` method for that fitted meta-learner.

What coefficients represent. These coefficients correspond to the meta-learner regression of the outcome on the cross-validated base-learner predictions (Z_1, Z_2, \dots). Under the default meta-learner, they are the stacking weights (on the scale defined by the meta-learner).

Learner-dependent output. The returned coefficient object depends on the meta-learner implementation (by default a `glmnet` fit, often returning a sparse matrix). This method does not rename Z^* terms or post-process coefficients; it returns the output of `coef(object$superlearner[[k]]$meta_learner_fit, ...)` unchanged.

Single-learner special case. If the ensemble was fit with only one base learner, no meta-learner is fit and `meta_learner_fit` is NULL. In that case, `coef()` for the `poisson_superlearner` does not have meta-learner coefficients to return.

Value

If `cause` is a single integer, returns the coefficient object produced by `coef()` for the selected cause-specific fitted model: the meta-learner when `model = "s1"` and a meta-learner is available, or the selected base learner when `model` selects a base learner or no meta-learner is available.

If `cause = NULL`, returns a list of length `object$data_info$n_crisis`, where element `[[k]]` contains coefficients for the selected model for cause k .

If no fitted ensemble is present (`object$superlearner` is NULL), signals a message and returns `invisible(object)`.

Examples

```
d <- simulateStenoT1(50, competing_risks = TRUE)
learners <- list(
  glm = Learner_glmnet(covariates = c("age", "value_LDL"), lambda = 0, cross_validation = FALSE),
  gam = Learner_gam(covariates = c("age", "value_LDL"))
)
fit <- Superlearner(d, id="id", status="status_cvd", event_time="time_cvd",
  learners=learners, number_of_nodes=4, nfold=2)

# meta-learner coefficients (cause 1)
coef(fit, cause = 1)

# meta-learner coefficients for all causes (list)
coef(fit)
```

fit_learner

Fit a single base learner

Description

Pre-processes subject-level time-to-event data into a long Poisson format on a piecewise-constant time grid, then fits **one** initialized learner object. For competing risks, a separate model is fit for each event type (cause) using the standard cause-specific Poisson likelihood on the long data.

Usage

```
fit_learner(
  data,
  learner,
  id = "id",
  stratified_k_fold = FALSE,
  status = "status",
  event_time = NULL,
  number_of_nodes = NULL,
  nodes = NULL,
  variable_transformation = NULL,
  ...
)
```

Arguments

data	data.frame. Subject-level input data (one row per subject).
learner	Reference-class learner object (e.g. from Learner_glmnet() , Learner_hal() or Learner_gam()). Must implement a <code>\$private_fit(dt_long)</code> method that fits the learner on long Poisson data for one cause.
id	character(1). Name of the subject identifier column. If not found in data, an id column is created automatically.

stratified_k_fold	logical(1). Reserved argument for future fold strategy. Currently ignored.
status	character(1). Name of the event-status column. Must be coded with 0 = censoring and 1, 2, ..., K for event types (causes). If there is no 0 in status, the data are treated as uncensored.
event_time	character(1). Name of the event/censoring time column. Must be present in data.
number_of_nodes	numeric(1) or NULL. If not NULL, constructs a quantile-based node grid with number_of_nodes + 1 cut points (including endpoints), then adds 0 if missing.
nodes	numeric or NULL. Explicit time-node grid (cut points). If supplied, number_of_nodes is ignored. 0 is added if missing. Nodes beyond max(event_time) are dropped.
variable_transformation	list/character/formula or NULL. Optional transformations applied to the internally created long Poisson data before fitting (via apply_transformations()).
...	Additional arguments currently ignored.

Value

An object of class `base_learner`, i.e. a named list with:

model The **learner object** that was fit (the input learner), stored for later prediction. This contains the learner specification (e.g., covariates, tuning parameters).

learner_fit A list of fitted model objects, **one per cause**. Its length equals `data_info$n_crisks`. The list is created by splitting the internally pre-processed long data by cause indicator `k` and calling `model$private_fit()` on each split.

- Names typically correspond to the cause labels "1", "2", ..., "K".
- Each element is **learner-dependent**: e.g. for `Learner_glmnet` it may be a "glmnet" (often wrapped, e.g. "fishnet") fit; for other learners it will be whatever `$private_fit()` returns.
- Each fitted object is trained on long Poisson data representing the piecewise-constant hazard for that cause across the node intervals.

data_info A list of bookkeeping information needed for prediction and interpretation:

id Identifier column name used.

status Status column name used.

event_time Event/censoring time column name used.

nodes Numeric vector of node cut points used for the piecewise grid (includes 0 and is sorted). These are the interval boundaries used in the long Poisson representation.

maximum_followup `max(data[[event_time]])`.

n_crisks Number of event types (causes) detected. If censoring is present (0 in status), then `n_crisks = #unique(status) - 1`; otherwise `n_crisks = #unique(status)`.

variable_transformation The transformation specification passed in `variable_transformation` (or NULL).

Examples

```
d <- simulateStenoT1(50, competing_risks = TRUE)
lrn <- Learner_glmnet(covariates = c("age", "value_LDL"),
                     lambda = 0,
                     cross_validation = FALSE)
bl <- fit_learner(d,
                 learner = lrn,
                 id = "id",
                 status = "status_cvd",
                 event_time = "time_cvd",
                 number_of_nodes = 2)
```

Learner_gam-class	GAM learner via mgcv::bam
-------------------	---------------------------

Description

Learner_gam is a Reference Class implementing the learner interface used by [Superlearner\(\)](#) and [fit_learner\(\)](#).

Arguments

covariates character. Right-hand-side terms, including mgcv smooths (e.g. "s(age)") and/or linear terms (e.g. "value_LDL").

cross_validation logical. Included for compatibility with the learner interface; smoothing selection is controlled by mgcv and arguments in

Details

User-facing API: users should **only initialize** the learner and pass it to [Superlearner\(\)](#) / [fit_learner\(\)](#). The remaining methods documented below are part of the internal learner interface and are **not meant to be called directly by users**.

Wrapper role: this class wraps `mgcv::bam` in a piecewise-constant hazard workflow. The package-specific contribution is to provide a convenient interface for the long-format Poisson likelihood with offsets for time at risk, and optional node terms encoding the baseline hazard, while forwarding standard `mgcv::bam` arguments supplied via

Model

Let $0 = t_0 < t_1 < \dots < t_m$ denote time knots and define interval indicators $I_k(t) = 1\{t \in (t_k, t_{k+1}]\}$. The piecewise-constant hazard model with an additive predictor is

$$\lambda(t | x) = \sum_{k=0}^m I_k(t) \exp\{\eta(x) + \gamma_k\}.$$

The additive predictor $\eta(x)$ is constructed from `covariates` (smooth terms such as `s(age)` and/or linear terms) and estimated by mgcv.

Fields

`covariates` (character) Terms used to build the additive predictor (may include `s()` terms).
`cross_validation` (logical) Workflow flag; see Details.
`intercept` (logical) Whether to include an intercept.
`formula` (character) Formula string passed to `mgcv::bam`.
`learner` (function) Backend fitter (`mgcv::bam`).
`fit_arguments` (list) Additional arguments forwarded to `mgcv::bam`.

Methods (internal learner interface)

`initialize(...)` Construct and configure the learner. This is the only method users should call.
`private_fit(data, ...)` Internal. Fits a Poisson GAM with offset $\log(t_{ij})$ on long-format data.
`private_fit_all_causes(data, ...)` Internal. Fits cause-specific Poisson GAMs for all requested causes using a shared long-format data setup.
`private_predictor(model, newdata, ...)` Internal. Predicts hazards on the response scale.

Examples

```
lrm <- Learner_gam(covariates = c("s(age)", "value_LDL"))
```

Learner_glmnet-class *Penalized Poisson learner via glmnet*

Description

Learner_glmnet is a Reference Class implementing the learner interface used by [Superlearner\(\)](#) and [fit_learner\(\)](#).

Details

User-facing API: users are expected to **initialize** the learner (i.e., call `Learner_glmnet(...)`) and pass the resulting object to [Superlearner\(\)](#) or [fit_learner\(\)](#). The remaining methods documented below are part of the internal learner interface and are **not meant to be called directly by users**.

Wrapper role: this class is a user-friendly wrapper around the existing `glmnet` implementation. The package-specific contribution is to provide a piecewise-constant hazard workflow: create the long-format Poisson data with offsets for time at risk, include interval ("node") indicators for the baseline hazard, and forward standard `glmnet` arguments supplied at initialization to the backend fitter.

Model

Let $0 = t_0 < t_1 < \dots < t_m$ denote time knots and define interval indicators $I_k(t) = 1\{t \in (t_k, t_{k+1}]\}$. The piecewise-constant hazard model is

$$\lambda(t | x) = \sum_{k=0}^m I_k(t) \lambda_k(x), \quad \lambda_k(x) = \exp(\beta^\top x + \gamma_k).$$

Penalization is applied to the regression coefficients through the glmnet elastic-net penalty. Node (baseline) terms are given zero penalty by default; if this backend call fails, the learner retries with a fully penalized design.

Fields

- covariates (character) Names of covariate columns used in the model.
- cross_validation (logical) If TRUE, chooses lambda by `glmnet::cv.glmnet`.
- intercept (logical) Backend intercept flag; currently fixed to TRUE by the constructor.
- lambda (numeric) If `cross_validation=FALSE`, the lambda used in the final fit.
- formula (character) Formula string used to create the design matrix in long format.
- learner (function) Backend fitter (`glmnet::glmnet` or `glmnet::cv.glmnet`).
- fit_arguments (list) Additional arguments forwarded to the backend fitter.

Methods (internal learner interface)

- `initialize(...)` Construct and configure the learner. This is the only method users should call.
- `private_fit(data, ...)` Internal. Fits a Poisson model with offset `log(tij)` on long-format data.
- `private_fit_all_causes(data, ...)` Internal. Fits cause-specific Poisson models for all requested causes using a shared long-format data setup.
- `private_predictor(model, newdata, ...)` Internal. Predicts hazards on the response scale for long-format newdata.

Examples

```
lrn <- Learner_glmnet(covariates = c("age", "sex"), alpha = 1, cross_validation = TRUE)
```

Learner_hal-class *HAL learner for piecewise Poisson hazards*

Description

Learner_hal is a Reference Class implementing the learner interface used by [Superlearner\(\)](#) and [fit_learner\(\)](#).

Details

User-facing API: users should **only initialize** the learner and pass it to `Superlearner()/fit_learner()`. The remaining methods documented below are part of the internal learner interface and are **not meant to be called directly by users**.

Wrapper role: this class provides a piecewise-constant hazard wrapper around a HAL-style indicator-basis construction, estimated by L1-penalized Poisson regression using a `glmnet` backend. The package-specific contribution is to (i) construct the long-format Poisson representation with offsets for time at risk, (ii) generate indicator bases compatible with piecewise hazards, and (iii) forward backend fitting arguments supplied via

Model

Let $0 = t_0 < t_1 < \dots < t_m$ denote time knots and define interval indicators $I_k(t) = 1\{t \in (t_k, t_{k+1}]\}$. The HAL piecewise-constant hazard model is

$$\lambda(t | x) = \sum_{k=0}^m I_k(t) \exp\{f(t, x)\},$$

where $f(t, x)$ is approximated by a finite linear combination of indicator basis functions.

Two-covariate illustration

Let $x = (x_1, x_2)$ be two covariates and let $t_0 < t_1 < \dots < t_R$ be time grid points used to create step functions in time. Choose covariate cutpoints $c_{1,1}, \dots, c_{1,K_1}$ for x_1 and $c_{2,1}, \dots, c_{2,K_2}$ for x_2 .

Define indicator bases:

$$\begin{aligned} B_r(t) &= 1\{t_r \leq t\} \\ B_{1,p}(x) &= 1\{c_{1,p} \leq x_1\} \\ B_{2,q}(x) &= 1\{c_{2,q} \leq x_2\} \end{aligned}$$

A main-effects HAL approximation on the log-hazard scale can be written as:

$$f_\beta(t, x) = \beta_0 + \sum_{r=1}^R \beta_r B_r(t) + \sum_{r=1}^R \sum_{p=1}^{K_1} \beta_{r,1,p} B_r(t) B_{1,p}(x) + \sum_{r=1}^R \sum_{q=1}^{K_2} \beta_{r,2,q} B_r(t) B_{2,q}(x).$$

If `max_degree` ≥ 2 , the learner additionally includes interaction bases such as

$$\sum_{r=1}^R \sum_{p=1}^{K_1} \sum_{q=1}^{K_2} \beta_{r,12,pq} B_r(t) B_{1,p}(x) B_{2,q}(x).$$

How reference class parameters map to the model

`covariates` Covariate columns used to build covariate indicator bases.

`num_knots` Controls the number of cutpoints per covariate used for indicator bases.

`max_degree` Maximum interaction order included in the basis expansion.

`intercept` Whether the backend penalized regression includes an intercept term.

`cross_validation` If TRUE, selects the penalty level using `glmnet::cv.glmnet`.
`maxit_prefit` Optional `maxit` value used for the initial HAL backend fit. Leave as NA to use the backend default.
`fit_arguments` Additional arguments forwarded to the `glmnet` backend (e.g. `nfolds`).

Fields

`covariates` (character) Names of covariate columns used in the basis.
`cross_validation` (logical) Whether to use `cv.glmnet` to select the penalty.
`intercept` (logical) Backend intercept flag.
`max_degree` (integer) Maximum interaction order.
`num_knots` (numeric) Knots used for basis construction.
`lambda_opt` (numeric) Selected penalty level when using cross-validation.
`maxit_prefit` (numeric) Optional `maxit` value used for the initial HAL backend fit.
`fit_arguments` (list) Extra backend arguments forwarded to `glmnet`.

Methods (internal learner interface)

`initialize(...)` Construct and configure the learner. This is the only method users should call.
`hal_basis(...)` Internal helper. Constructs HAL basis matrices and metadata for fitting.
`hal_prepare_new(...)` Internal helper. Builds prediction-time HAL basis matrices from fitted basis metadata.
`private_fit(data, ...)` Internal. Builds bases and fits the penalized Poisson model with offset $\log(t_{ij})$.
`private_fit_all_causes(data, ...)` Internal. Fits penalized Poisson HAL models for all requested causes using a shared basis setup.
`private_predictor(model, newdata, ...)` Internal. Evaluates the fitted approximation and returns hazards on the response scale.

Examples

```
lrn <- Learner_hal(covariates = c("age", "sex"), max_degree = 2L, num_knots = c(10L, 5L))
```

<code>pch_absolute_risk</code>	<i>Absolute risk (cumulative incidence) for a cause under piecewise-constant hazards</i>
--------------------------------	--

Description

Computes, per row, the cumulative incidence function at the end of each interval, grouped by `id`. The number of causes is inferred from the number of columns in `haz`.

Usage

```
pch_absolute_risk(id, dt, haz, cause_idx, one_based = TRUE, na_is_zero = FALSE)
```

Arguments

id	Integer vector. Sorted by id then time.
dt	Numeric vector of interval lengths.
haz	Numeric matrix (n x C) of cause-specific hazards per interval. Columns correspond to causes 1..C.
cause_idx	Integer. Index of the cause of interest (1-based by default).
one_based	Logical. If TRUE, cause_idx is 1-based. If FALSE, 0-based.
na_is_zero	Logical. If TRUE, treat NA/Inf hazards as zero.

Value

Numeric vector of cumulative incidence values at the end of each interval.

Examples

```
id <- c(1L, 1L, 2L, 2L)
dt <- c(1, 1, 1, 1)
haz <- rbind(
  c(0.10, 0.05),
  c(0.20, 0.10),
  c(0.05, 0.02),
  c(0.10, 0.03)
)
pch_absolute_risk(id = id, dt = dt, haz = haz, cause_idx = 1)
```

pch_absolute_risk_euler

Absolute risk (Euler approximation) for a cause under piecewise-constant hazards

Description

Computes the cumulative incidence function using the first-order Euler (discrete) approximation:

$$F_j(t) \approx \sum S(t_{k-1}) \lambda_{j,k} \Delta t_k$$

Grouped by id, this returns the cumulative incidence at the end of each interval.

Usage

```
pch_absolute_risk_euler(
  id,
  dt,
  haz,
  cause_idx,
  one_based = TRUE,
  na_is_zero = FALSE
)
```

Arguments

id	Integer vector. Sorted by id then time.
dt	Numeric vector of interval lengths.
haz	Numeric matrix (n x C) of cause-specific hazards per interval.
cause_idx	Integer. Index of the cause of interest (1-based by default).
one_based	Logical. If TRUE, cause_idx is 1-based. If FALSE, 0-based.
na_is_zero	Logical. If TRUE, treat NA/Inf hazards as zero.

Value

Numeric vector of cumulative incidence values (Euler approximation) at the end of each interval.

Examples

```
id <- c(1L, 1L, 2L, 2L)
dt <- c(1, 1, 1, 1)
haz <- rbind(
  c(0.10, 0.05),
  c(0.20, 0.10),
  c(0.05, 0.02),
  c(0.10, 0.03)
)
pch_absolute_risk_euler(id = id, dt = dt, haz = haz, cause_idx = 1)
```

pch_survival

Piecewise-constant hazards survival function

Description

Computes survival at the end of each interval for competing risks with piecewise constant hazards.

Usage

```
pch_survival(id, dt, haz, na_is_zero = FALSE)
```

Arguments

id	Integer vector of subject IDs, sorted by id then time.
dt	Numeric vector of interval lengths.
haz	Numeric matrix (n x C) of cause-specific hazards.
na_is_zero	Logical. If TRUE, treat NA hazards as zero.

Value

Numeric vector of survival probabilities at the end of each interval.

Examples

```
id <- c(1L, 1L, 2L, 2L)
dt <- c(1, 1, 1, 1)
haz <- rbind(
  c(0.10, 0.05),
  c(0.20, 0.10),
  c(0.05, 0.02),
  c(0.10, 0.03)
)
pch_survival(id = id, dt = dt, haz = haz)
```

predict.base_learner *Predict hazards, survival and absolute risk from a fitted base learner*

Description

Computes **cause-specific piecewise-constant hazards** (pwch_k), the corresponding **survival function**, and **absolute risk** for a given cause, at user-supplied prediction horizons times, using a fitted base_learner object (single learner; no stacking).

Usage

```
## S3 method for class 'base_learner'
predict(object, newdata, times, cause = 1, ...)
```

Arguments

object	base_learner. A fitted object returned by <code>fit_learner()</code> . It contains the learner specification in <code>object\$model</code> and cause-specific fitted models in <code>object\$learner_fit</code> .
newdata	data.frame/data.table. New covariate data (one row per subject). If newdata contains the original event_time, status, or id columns used for fitting, they are ignored for prediction.

times	numeric. Prediction horizon(s). May include 0. Times larger than <code>object\$data_info\$maximum_follow</code> are not supported: if all requested times exceed the maximum follow-up, a warning is issued and NULL is returned; if only some exceed, output rows for those times are returned with NA predictions.
cause	numeric(1). Cause index (1, 2, ...) used for the <code>absolute_risk</code> calculation.
...	Additional arguments (currently ignored).

Details

Internally, `newdata` is expanded to a Cartesian product with `times`, converted to long Poisson format on `object$data_info$nodes`, and the fitted learner for each cause in `object$learner_fit` is used to predict the cause-specific hazards. Survival and absolute risk are then computed from the predicted hazards.

Special case `times = 0`: when 0 is included in `times`, the returned rows have `survival_function = 1`, `absolute_risk = 0`, and all `pwch_k = 0` at time 0.

Identifiers in the output: if `newdata` contains the `id` column, it is carried into the output. If `newdata` does not contain an `id` column, an internal `id` is created for computation, but it is not guaranteed to appear in the returned table unless it was present in `newdata`.

Value

A `data.table` with one row per (row in `newdata`, time in `times`) and columns:

(original columns) All columns from `newdata` (excluding ignored event columns).

time column A column with name `object$data_info$event_time` holding the requested horizon.

pwch_1, pwch_2, ... Predicted cause-specific piecewise hazards at the horizon.

survival_function Predicted survival probability at the horizon.

absolute_risk Predicted cumulative incidence (absolute risk) for cause at the horizon.

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)
lrn <- Learner_glmnet(covariates = c("age", "value_LDL"), lambda = 0, cross_validation = FALSE)
bl <- fit_learner(d, learner = lrn, id="id", status="status_cvd", event_time="time_cvd",
                 number_of_nodes=8)
p <- predict(bl, newdata = d[1:5], times = c(0, 2, 5), cause = 1)
head(p)
```

```
predict.poisson_superlearner
```

Predict hazards, survival and absolute risk from a fitted Poisson Super Learner

Description

Computes **cause-specific piecewise-constant hazards** (pwch_k), the corresponding **survival function**, and **absolute risk** for a given cause, at user-supplied prediction horizons `times`, for each row in `newdata`.

Usage

```
## S3 method for class 'poisson_superlearner'
predict(object, newdata, times, cause = 1, model = "sl", ...)
```

Arguments

<code>object</code>	<code>poisson_superlearner</code> . A fitted ensemble from Superlearner() .
<code>newdata</code>	<code>data.frame/data.table</code> . New covariate data (one row per subject). If <code>newdata</code> contains the original <code>event_time</code> , <code>status</code> , or <code>id</code> columns used for fitting, they are ignored for prediction.
<code>times</code>	numeric. Prediction horizon(s). May include 0. Times larger than <code>object\$data_info\$maximum_follow</code> are not supported: if all requested times exceed the maximum follow-up, a warning is issued and NULL is returned; if only some exceed, output rows for those times are returned with NA predictions.
<code>cause</code>	numeric(1). Cause index (1, 2, ...) used for the <code>absolute_risk</code> calculation.
<code>model</code>	Model selector. Default is "sl" for the stacked super learner. Allowed values are: 0, "sl", "superlearner", or "super_learner" Use the stacked super learner prediction. For causes with only one retained learner or no fitted meta-learner, this falls back to the retained base learner for that cause. "discrete_sl" and aliases For each cause, use the retained base learner with the smallest cross-validated deviance. learner label Use one stored base learner by its label in <code>object\$data_info\$learners_labels[[k]]</code> . "learner_j" or character integer "j" Use the j-th stored learner. integer j >= 1 Use the j-th stored learner. vector of labels or positive integer indices Use cause-specific base learners; length must equal <code>object\$data_info\$n_crisks</code> . Numeric positions refer to the learners actually retained for each cause in the fitted object.
<code>...</code>	Additional arguments (currently ignored).

Details

Internally, newdata is expanded to a Cartesian product with the requested times, converted to long Poisson format on `object$data_info$nodes`, and hazards are predicted either from the stacked super learner (`model = "sl"`), the discrete super learner (`model = "discrete_sl"`), or selected fitted base learners. Survival and absolute risk are then computed from the predicted hazards.

Special case `times = 0`: when 0 is included in times, the returned rows have `survival_function = 1`, `absolute_risk = 0`, and all `pwch_k = 0` at time 0.

Identifiers in the output: if newdata contains the `id` column, it is carried into the output. If newdata does not contain an `id` column, an internal `id` is created for computation, but it is not guaranteed to appear in the returned table unless it was present in newdata.

Value

A `data.table` with one row per (row in newdata, time in times) and columns:

(original columns) All columns from newdata (excluding ignored event columns).

time column A column with name `object$data_info$event_time` holding the requested horizon.

pwch_1, pwch_2, ... Predicted cause-specific piecewise hazards at the horizon.

survival_function Predicted survival probability at the horizon.

absolute_risk Predicted cumulative incidence (absolute risk) for cause at the horizon.

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)
```

```
learners <- list(
  lasso = Learner_glmnet(
    covariates = "sex",
    alpha = 1,
    lambda = 0.01,
    cross_validation = FALSE
  ),
  ridge = Learner_glmnet(
    covariates = c("sex", "value_LDL"),
    alpha = 0,
    lambda = 0.01,
    cross_validation = FALSE
  )
)
```

```
fit <- Superlearner(
  data = d,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  learners = learners,
  number_of_nodes = 3,
  nfold = 2
)
```

```

)
p <- predict(fit, newdata = d[1:3], times = c(0, 2), cause = 1)
p[, .(id, time_cvd, absolute_risk)]

```

predictRisk.base_learner

Absolute-risk matrix predictions for a fitted base learner

Description

Absolute-risk matrix predictions for a fitted base learner

Usage

```

## S3 method for class 'base_learner'
predictRisk(object, newdata, times, cause = 1, ...)

```

Arguments

object	base_learner. Fitted object from <code>fit_learner()</code> .
newdata	data.frame. New covariate data.
times	numeric. Prediction times.
cause	numeric(1). Cause index.
...	Unused.

Value

numeric matrix with `nrow(newdata)` rows and `length(times)` columns.

Examples

```

d <- simulateStenoT1(30, competing_risks = TRUE)
lrn <- Learner_glmnet(
  covariates = c("sex", "value_LDL"),
  lambda = 0.01,
  cross_validation = FALSE
)
bl <- fit_learner(
  d,
  learner = lrn,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  number_of_nodes = 3
)

```

```

if (requireNamespace("riskRegression", quietly = TRUE)) {
  riskRegression::predictRisk(bl, newdata = d[1:3], times = c(1, 3), cause = 1)
}

```

predictRisk.poisson_superlearner

Absolute-risk matrix predictions for a fitted Poisson Super Learner

Description

S3 method compatible with `riskRegression::predictRisk` returning one column per requested time.

Usage

```

## S3 method for class 'poisson_superlearner'
predictRisk(object, newdata, times, cause = 1, model = "s1", ...)

```

Arguments

<code>object</code>	poisson_superlearner. Fitted object.
<code>newdata</code>	data.frame. New covariate data.
<code>times</code>	numeric. Prediction times.
<code>cause</code>	numeric(1). Cause index.
<code>model</code>	Model selector. Default is "s1". Allowed values are the same as in predict.poisson_superlearner() , including "discrete_s1" and cause-specific vectors of base-learner labels or indices.
<code>...</code>	Unused.

Value

numeric matrix with `nrow(newdata)` rows and `length(times)` columns.

Examples

```

d <- simulateStenoT1(30, competing_risks = TRUE)

learners <- list(
  lasso = Learner_glmnet(
    covariates = "sex",
    alpha = 1,
    lambda = 0.01,
    cross_validation = FALSE
  ),
  ridge = Learner_glmnet(

```

```

    covariates = c("sex", "value_LDL"),
    alpha = 0,
    lambda = 0.01,
    cross_validation = FALSE
  )
)

fit <- Superlearner(
  data = d,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  learners = learners,
  number_of_nodes = 3,
  nfold = 2
)

if (requireNamespace("riskRegression", quietly = TRUE)) {
  riskRegression::predictRisk(fit, newdata = d[1:3], times = c(1, 3), cause = 1)
}

```

```
print.base_learner    Print method for base_learner
```

Description

Prints a compact description of the fitted base learner, including the learner type, the time-grid used, and (optionally) the fitted model object for a given cause.

Usage

```
## S3 method for class 'base_learner'
print(x, cause = 1, ...)
```

Arguments

x	base_learner object returned by <code>fit_learner()</code> .
cause	numeric(1) or NULL. Which cause to print the fitted model for. If NULL, prints one line per cause (classes only) instead of printing the full fitted objects.
...	Passed to the underlying fitted object <code>print()</code> method when cause is a single integer.

Value

Invisibly returns x.

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)
lrn <- Learner_glmnet(
  covariates = c("sex", "value_LDL"),
  lambda = 0.01,
  cross_validation = FALSE
)
bl <- fit_learner(
  d,
  learner = lrn,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  number_of_nodes = 3
)
print(bl, cause = NULL)
```

```
print.poisson_superlearner
```

Print method for poisson_superlearner

Description

Prints a compact description of the fitted Poisson Super Learner, including the number of base learners, the meta-learner, the time-grid used, and competing-risk structure. Optionally prints the fitted meta-learner for a given cause.

Usage

```
## S3 method for class 'poisson_superlearner'
print(x, cause = 1, model = "sl", ...)
```

Arguments

x	poisson_superlearner object returned by Superlearner() .
cause	numeric(1) or NULL. Which cause's meta-learner fit to print. If NULL, prints one line per cause (classes only) instead of printing the full fitted objects.
model	Model selector. Default is "sl" for the stacked super learner. Allowed values are: 0, "sl", "superlearner", or "super_learner" Print the stacked meta-learner. For causes with no fitted meta-learner, this falls back to the retained base learner. "discrete_sl" and aliases Print the cause-specific base learner with the smallest cross-validated deviance. learner label Print one stored base learner by its label in x\$data_info\$learners_labels[[k]].

"learner_j" or character integer "j" Print the j-th stored learner.
integer j >= 1 Print the j-th stored learner.
vector of labels or positive integer indices Use cause-specific base learners; length must equal x\$data_info\$n_crisks.
 ... Passed to the underlying fitted meta-learner print() method when cause is a single integer.

Value

Invisibly returns x.

Examples

```

d <- simulateStenoT1(30, competing_risks = TRUE)

learners <- list(
  lasso = Learner_glmnet(
    covariates = "sex",
    alpha = 1,
    lambda = 0.01,
    cross_validation = FALSE
  ),
  ridge = Learner_glmnet(
    covariates = c("sex", "value_LDL"),
    alpha = 0,
    lambda = 0.01,
    cross_validation = FALSE
  )
)

fit <- Superlearner(
  data = d,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  learners = learners,
  number_of_nodes = 3,
  nfold = 2
)

print(fit, cause = NULL)

```

simulateStenoT1

Simulate time-to-event data for hypothetical type-1 diabetes patients

Description

Simulate synthetic data inspired by the Steno Type-1 risk engine

Usage

```
simulateStenoT1(
  n,
  coefficient_age = 0.05,
  coefficient_LDL = 0.1,
  value_diabetis = 0.02,
  seed = NULL,
  keep = NULL,
  scenario = c("alpha", "beta"),
  competing_risks = FALSE
)
```

Arguments

<code>n</code>	numeric(1). Number of subjects to simulate.
<code>coefficient_age</code>	numeric(1). Log-hazard coefficient for age in the CVD model (time.event.1).
<code>coefficient_LDL</code>	numeric(1). Log-hazard coefficient for LDL in the CVD model (time.event.1).
<code>value_diabetis</code>	numeric(1). Log-hazard coefficient for diabetes duration in the CVD model (time.event.1).
<code>seed</code>	integer(1) or NULL. Optional random seed passed to <code>set.seed()</code> before simulating the data. If NULL, the current RNG state is used.
<code>keep</code>	character or NULL. Optional subset of columns to retain. If supplied, only those columns are returned.
<code>scenario</code>	character(1). One of "alpha" or "beta". Scenario "beta" modifies the CVD hazard by adding nonlinear hinge-squared terms in age and LDL.
<code>competing_risks</code>	logical(1). If TRUE, simulates two event causes (CVD and death without CVD). Otherwise simulates CVD vs censoring.

Details

Generates baseline covariates and event times for CVD and censoring, with an optional competing-risks setting, for examples, benchmarks and tests.

The simulator uses a structural equation model (via `lava : lvm`) to generate realistic correlations between covariates. Event times are then generated from cause-specific Weibull proportional hazards models, where the linear predictor depends on the simulated covariates (and scenario).

The following baseline covariates are generated (column name, type, interpretation):

sex factor. Binary sex indicator (generated Bernoulli, then stored as factor).

age numeric. Age at baseline (years).

diabetes_duration numeric. Duration of diabetes at baseline (years).

value_SBP numeric. Systolic blood pressure (SBP).

value_LDL numeric. LDL cholesterol.

value_HbA1C numeric. HbA1c.

value_Albuminuria factor with levels Normal, Micro, Macro. Albuminuria category.

eGFR numeric. Estimated glomerular filtration rate, constructed from latent age-dependent log2 eGFR components (higher values indicate better kidney function).

value_Smoking factor. Smoking indicator (generated from a logistic model, then stored as factor).

value_Motion factor. Physical activity indicator (generated from a logistic model, then stored as factor).

Event time variables are generated from latent Weibull PH models: `time.event.1` (CVD), `time.event.0` (censoring), and, if `competing_risks = TRUE`, `time.event.2` (death without prior CVD). These latent variables are used to construct the observed outcome variables returned by the function (see below).

Value

A data table with at least the following columns:

id integer. Subject identifier (1, ..., n).

time_cvd numeric. Observed follow-up time (minimum of event and censoring times; also includes competing risk time if `competing_risks = TRUE` in scenario "alpha").

status_cvd integer. Observed event status: 0 = censored, 1 = CVD, and if `competing_risks = TRUE` in scenario "alpha", 2 = death without prior CVD.

time numeric. Alias of `time_cvd` (kept for convenience).

event integer. Alias of `status_cvd` (kept for convenience).

uncensored_time_cvd numeric. Event time ignoring censoring (minimum of event causes only).

uncensored_status_cvd integer. Event cause ignoring censoring. In scenario "alpha" this is 1 (CVD) or 2 (death without CVD); in scenario "beta" this is always 1.

uncensored_time numeric. Alias of `uncensored_time_cvd`.

uncensored_event integer. Alias of `uncensored_status_cvd`.

In addition, the returned table contains all baseline covariates listed in **Details**. Internal latent variables used only for simulation are removed before returning (e.g., log2 eGFR components and, in scenario "beta", the hinge-squared features).

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Examples

```
simulateStenoT1(n = 20, scenario = "alpha", competing_risks = TRUE)
```

summary.base_learner *Summarize a fitted base learner object*

Description

Dispatches to the underlying fitted model's `summary()` method for the selected cause, or returns a list of summaries for all causes.

Usage

```
## S3 method for class 'base_learner'  
summary(object, cause = 1, ...)
```

Arguments

<code>object</code>	base_learner returned by <code>fit_learner()</code> .
<code>cause</code>	numeric(1) or NULL. Which cause to summarize. If NULL, returns one summary per cause.
<code>...</code>	Passed to the underlying <code>summary()</code> method (learner-dependent).

Value

If `cause` is a single integer, returns the underlying model summary for that cause. If `cause = NULL`, returns a list of summaries (one per cause).

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)  
lrn <- Learner_glmnet(  
  covariates = c("sex", "value_LDL"),  
  lambda = 0.01,  
  cross_validation = FALSE  
)  
bl <- fit_learner(  
  d,  
  learner = lrn,  
  id = "id",  
  status = "status_cvd",  
  event_time = "time_cvd",  
  number_of_nodes = 3  
)  
out <- summary(bl, cause = 1)
```

```
summary.poisson_superlearner
```

Summarize a fitted Poisson Super Learner object

Description

Prints:

1. a compact description of the fitted ensemble,
2. cross-validated deviances for base learners (when available),
3. cause-specific meta-learner coefficients (stacking weights).

Usage

```
## S3 method for class 'poisson_superlearner'
summary(object, cause = NULL, model = "sl", ...)
```

Arguments

object	poisson_superlearner returned by Superlearner() .
cause	numeric(1) or NULL. Which cause's meta-learner fit to print. If NULL, prints one line per cause (classes only) instead of printing the full fitted objects.
model	Model selector. Default is "sl" for the stacked super learner. Allowed values are: 0, "sl", "superlearner", or "super_learner" Summarize the stacked meta-learner. For causes with no fitted meta-learner, this falls back to the retained base learner. "discrete_sl" and aliases Summarize the cause-specific base learners with the smallest cross-validated deviance. learner label Summarize one stored base learner by its label in object\$data_info\$learners_labels["learner_j" or character integer "j" Summarize the j-th stored learner. integer j >= 1 Summarize the j-th stored learner. vector of labels or positive integer indices Use cause-specific base learners; length must equal object\$data_info\$n_crisks.
...	Passed to the underlying coef() method for the fitted meta-learner (learner-dependent; e.g. s for glmnet).

Value

Invisibly returns a list with elements:

cross_validation_deviance data.table (or NULL).

meta_coefficients List of length n_crisks with cause-specific coefficient objects (or NULL).

Examples

```
d <- simulateStenoT1(30, competing_risks = TRUE)

learners <- list(
  lasso = Learner_glmnet(
    covariates = "sex",
    alpha = 1,
    lambda = 0.01,
    cross_validation = FALSE
  ),
  ridge = Learner_glmnet(
    covariates = c("sex", "value_LDL"),
    alpha = 0,
    lambda = 0.01,
    cross_validation = FALSE
  )
)

fit <- Superlearner(
  data = d,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  learners = learners,
  number_of_nodes = 3,
  nfold = 2
)

s <- summary(fit, cause = 1)
names(s)
```

Superlearner

Fit a Poisson Super Learner ensemble

Description

Fits an ensemble of cause-specific piecewise-constant hazard models using a long-format Poisson representation and combines them through a meta-learner (stacking).

Usage

```
Superlearner(
  data,
  id = "id",
  status = "status",
  event_time = NULL,
  learners,
```

```

    number_of_nodes = NULL,
    nodes = NULL,
    variable_transformation = NULL,
    nfold = 3,
    verbose = FALSE,
    ...
  )

```

Arguments

<code>data</code>	<code>data.frame</code> . Subject-level input data, one row per subject.
<code>id</code>	<code>character(1)</code> . Name of the subject identifier column. If missing, an <code>id</code> column is created automatically.
<code>status</code>	<code>character(1)</code> . Name of the event-status column. It must be coded with <code>0</code> for censoring and <code>1, 2, ..., K</code> for event types. If there is no <code>0</code> in <code>status</code> , the data are treated as uncensored.
<code>event_time</code>	<code>character(1)</code> . Name of the event or censoring time column.
<code>learners</code>	<code>list</code> . Either a single learner library used for every cause, or a list of cause-specific learner libraries. A learner library is a named or unnamed list of initialized learner reference-class objects, for example <code>Learner_glmnet()</code> , <code>Learner_hal()</code> , or <code>Learner_gam()</code> . Missing learner names are filled as "learner_1", "learner_2", and so on; missing cause-library names are filled as "cause_1", "cause_2", and so on. Each learner must implement <code>\$private_fit(dt_long)</code> and <code>\$private_predictor(model, newdata)</code> .
<code>number_of_nodes</code>	<code>numeric(1)</code> or <code>NULL</code> . If not <code>NULL</code> , constructs a quantile-based node grid with <code>number_of_nodes + 1</code> cut points. Ignored when <code>nodes</code> is supplied.
<code>nodes</code>	<code>numeric</code> or <code>NULL</code> . Explicit time-node grid. If supplied, <code>number_of_nodes</code> is ignored. <code>0</code> is added if missing, and nodes larger than <code>max(event_time)</code> are dropped.
<code>variable_transformation</code>	Optional transformation specification passed to <code>apply_transformations()</code> on the internally created long-format data.
<code>nfold</code>	<code>numeric(1)</code> . Number of folds for cross-validation stacking.
<code>verbose</code>	<code>logical(1)</code> . If <code>TRUE</code> , display progress bars during full-data fitting and cross-validation fitting. Defaults to <code>FALSE</code> .
<code>...</code>	Additional arguments currently ignored.

Details

Internally, the function:

1. builds a time grid (`nodes`) and converts the subject-level data to a long Poisson format;
2. fits each base learner once on the full long data for each cause;
3. removes learners that already fail on the full data;
4. uses `nfold` cross-validation to obtain out-of-sample base-learner predictions (Z_1, Z_2, \dots) for stacking;

5. removes learners whose cross-validated prediction column is entirely missing for at least one cause;
6. fits a cause-specific meta-learner on the retained stacked predictions.

If all learners fail on the full data, the function stops with an error. If only one learner remains after the full-data screening step or after the cross-validation screening step, no meta-learner is fit. In that case, `metalearner` is `NULL`, each `superlearner[[k]]$meta_learner_fit` is `NULL`, and prediction is based directly on the stored fitted base learner. If some, but not all, causes retain only one learner after screening, those causes are predicted directly while other causes may still use a fitted meta-learner. Numeric learner positions always refer to the learners actually retained for the corresponding cause in the fitted object.

Value

An object of class `poisson_superlearner`, stored as a named list with the following components:

`learners`: a cause-specific list of retained base learner libraries. Thus `learners[[k]][[j]]` is the j -th retained learner object for cause k .

`metalearner`: a list describing the internal meta-learner used for stacking (`engine = "glmnet::glmnet"`, Poisson family, no intercept, `lambda = 0`, `add_nodes = FALSE`, log-hazard scale). If no stacking is performed because only one learner remains for every cause, `metalearner` is `NULL`.

`superlearner`: a list of length `data_info$n_crisks`, one entry per cause. For cause k , `superlearner[[k]]` is a list with two elements:

- `learners_fit`: the fitted base learner object or objects for cause k . If more than one learner is retained, this is a list with one fitted object per retained learner. If only one learner remains, this is the single fitted learner object itself.
- `meta_learner_fit`: the fitted cause-specific meta-learner for cause k . If no stacking is performed, this is `NULL`.

`cross_validation_deviance`: a `data.table` with columns `cause_index`, `cause`, `learner_index`, `learner`, and `deviance`, giving the cross-validated Poisson deviance for each retained base learner within each cause. This component is absent when all causes are fitted directly with a single retained learner.

`data_info`: a list of bookkeeping information used for prediction and interpretation, containing:

- `id`: identifier column name used.
- `status`: status column name used.
- `event_time`: event-time column name used.
- `nodes`: numeric vector of node cut points used for the piecewise grid.
- `nfold`: number of folds used for stacking.
- `maximum_followup`: maximum observed follow-up time.
- `n_crisks`: number of event types detected.
- `learners_labels`: list of character vectors with retained learner labels for each cause.
- `variable_transformation`: the transformation specification passed in `variable_transformation`, or `NULL`.

Examples

```
data <- simulateStenoT1(50, competing_risks = TRUE)

learners <- list(
  glm = Learner_glmnet(
    covariates = c("sex", "value_LDL"),
    lambda = 0,
    cross_validation = FALSE
  ),
  ridge = Learner_glmnet(
    covariates = c("sex", "value_LDL"),
    alpha = 0,
    lambda = 0.01,
    cross_validation = FALSE
  )
)

fit <- Superlearner(
  data = data,
  id = "id",
  status = "status_cvd",
  event_time = "time_cvd",
  learners = learners,
  number_of_nodes = 3,
  nfold = 2
)
```

Index

`coef.base_learner`, 2
`coef.poisson_superlearner`, 3

`fit_learner`, 5
`fit_learner()`, 2, 7–10, 14, 18, 20, 25

`Learner_gam` (`Learner_gam`-class), 7
`Learner_gam()`, 5, 28
`Learner_gam`-class, 7
`Learner_glmnet` (`Learner_glmnet`-class), 8
`Learner_glmnet()`, 5, 28
`Learner_glmnet`-class, 8
`Learner_hal` (`Learner_hal`-class), 9
`Learner_hal()`, 5, 28
`Learner_hal`-class, 9

`pch_absolute_risk`, 11
`pch_absolute_risk_euler`, 12
`pch_survival`, 13
`predict.base_learner`, 14
`predict.poisson_superlearner`, 16
`predict.poisson_superlearner()`, 19
`predictRisk.base_learner`, 18
`predictRisk.poisson_superlearner`, 19
`print.base_learner`, 20
`print.poisson_superlearner`, 21

`simulateStenoT1`, 22
`summary.base_learner`, 25
`summary.poisson_superlearner`, 26
`Superlearner`, 27
`Superlearner()`, 3, 4, 7–10, 16, 21, 26