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Title Bayesian Sample Size and Precision Considerations for Risk Prediction Models

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Description Performs Bayesian sample size, precision, and value-of-information analysis for external validation of existing multi-variable prediction models using the approach proposed by Sadatsafavi and colleagues (2026) <doi:10.1002/sim.70389>.

URL <https://github.com/resplab/bayespmtools>

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Contents

bpm_valprec	2
bpm_valsamp	4
calc_ciw_2s	6
calc_ciw_mc	6
calc_ciw_sample	7
calc_cstat	8
calc_riley_vars	8
calc_se_sp	9
find_n_mean	9
find_n_quantile	10
infer_cal_int_from_mean	11
infer_cal_int_from_oe	11
infer_correlation	12
inv_mean_quantile	12
inv_moments	13
isaric	13
moments	14
plot_cal_distance	14
plot_cal_instability	15
process_evidence	16
rbnorm	17
riley_samp	18
Index	19

bpm_valprec	<i>Bayesian Precision / VoI Calculator</i>
-------------	--

Description

Bayesian precision and value-of-information calculator for external validation studies of risk prediction models at fixed sample sizes.

Usage

```
bpm_valprec(
  N,
  evidence,
  targets,
  n_sim = NULL,
  method = "sample",
  threshold = NULL,
  dist_type = "logitnorm",
  impute_cor = TRUE,
  ex_args = NULL
)
```

Arguments

N	Numeric vector of sample sizes to evaluate.
evidence	A named list containing prior evidence components for model performance parameters (e.g., prevalence, discrimination, calibration). Alternatively, evidence may be a data frame of pre-posterior draws (element <code>\$sample</code>) returned by a previous call to this function or to <code>bpm_valprec()</code> , in which case those draws are used directly.
targets	A named list of targets to compute. ciw.metric Logical; compute expected CI width. qciw.metric Numeric scalar in (0,1); CI width quantile. oa.nb Logical; compute optimality assurance for net benefit. voi.nb Logical; compute EVSI and EVSI/EVPI.
n_sim	#' Number of Monte Carlo simulations used to generate the pre-posterior distribution. If evidence is a data frame from previous calls to relevant functions, n_sim will automatically be set to the number of rows of the data frame.
method	Method to compute CI widths. One of "sample" or "2s".
threshold	Decision threshold for net benefit calculations. Required if <code>oa.nb</code> or <code>voi.nb</code> are requested.
dist_type	Distribution for calibrated risks. Default is "logitnorm".
impute_cor	Logical; whether to induce correlation between parameters.
ex_args	Optional list of extra arguments. May include <code>f_progress</code> , a custom progress function.

Value

A list with elements:

results Matrix of requested metrics by sample size.

sample Monte Carlo sample used for computations.

evidence Processed evidence object.

targets Targets as supplied by the user.

ciws Simulated CI widths for requested metrics.

Examples

```
evidence <- list(
  prev ~ beta(116, 155),           # Outcome prevalence
  cstat ~ beta(3628, 1139),      # C-statistic
  cal_mean ~ norm(-0.009, 0.125), # Mean calibration error
  cal_slp ~ norm(0.995, 0.024)   # Calibration slope
)

res <- bpm_valprec(
  N = c(1000, 1500),
  evidence = evidence,
```

```

targets = list(eciw.cstat = TRUE, qciw.cal_slp=0.9, voi.nb=0.8),
threshold=0.2,
n_sim = 100      # faster and safer on CRAN. Please increase this value for real-world use.
)

print(res$results)

```

bpm_valsamp

Bayesian Sample Size Calculator for External Validation

Description

Bayesian sample size calculation for external validation studies of clinical risk prediction models. The function evaluates sample sizes required to meet precision-, assurance-, or decision-based targets using pre-posterior simulation.

Usage

```

bpm_valsamp(
  evidence,
  targets,
  n_sim = NULL,
  method = "sample",
  threshold = NULL,
  dist_type = "logitnorm",
  impute_cor = TRUE,
  ex_args = NULL
)

```

Arguments

- | | |
|----------|---|
| evidence | A named list containing prior evidence components for model performance parameters (e.g., prevalence, discrimination, calibration). Alternatively, evidence may be a data frame of pre-posterior draws (element <code>\$sample</code>) returned by a previous call to this function or to <code>bpm_valprec()</code> , in which case those draws are used directly. |
| targets | A named list specifying sample size targets.
Supported targets include: <ul style="list-style-type: none"> • Precision-based targets using expected 95\ (prefix <code>eciw</code>). • Assurance-based targets specifying the probability that the 95\ (prefix <code>qciw</code>). • Net benefit targets, including optimality assurance (<code>oa.nb</code>) and value-of-information ratios (<code>voi.nb = EVSI / EVPI</code>). |

For example, `eciw.cstat = 0.1` targets an expected interval width of 0.1 for the c-statistic, while `qciw.cal_slp = c(0.90, 0.22)` targets a 90 percent assurance that the calibration slope interval width does not exceed 0.22. Finally,

	<code>oa.nb = 0.80</code> targets a sample size that would correspond to 80 percent assurance that the strategy with the highest NB in the sample will be the strategy with the highest NB in the population.
<code>n_sim</code>	Number of Monte Carlo simulations used to generate the pre-posterior distribution. If evidence is a data frame from previous calls to relevant functions, <code>n_sim</code> will automatically be set to the number of rows of the data frame.
<code>method</code>	Method used to compute the pre-posterior distribution of 95\ One of "sample" (simulation-based) or "2s" (two-stage approximation). Default is "sample".
<code>threshold</code>	Risk threshold used for decision-analytic quantities and net benefit calculations. Required if <code>oa.nb</code> or <code>voi.nb</code> targets are specified.
<code>dist_type</code>	Distribution assumed for calibrated risks. Default is "logitnorm".
<code>impute_cor</code>	Logical indicating whether correlation between performance measures should be induced when simulating from marginal evidence distributions. Default is TRUE.
<code>ex_args</code>	Optional list of additional arguments passed to internal simulation or root-finding routines (experimental feature).

Value

A list with the following components:

- `results`: Estimated sample sizes required to meet each target.
- `sample`: Data frame of pre-posterior simulation draws.
- `evidence`: Processed evidence object used in the analysis.
- `trace`: Trace output from the stochastic root-finding algorithm.
- `targets`: The targets argument supplied to the function.

Examples

```
evidence <- list(
  prev ~ beta(116, 155),           # Outcome prevalence
  cstat ~ beta(3628, 1139),       # C-statistic
  cal_mean ~ norm(-0.009, 0.125), # Mean calibration error
  cal_slp ~ norm(0.995, 0.024)   # Calibration slope
)

targets <- list(
  eciw.cstat = 0.1,
  qciw.cstat = c(0.9, 0.1),
  oa.nb      = 0.8
)

samp <- bpm_valsamp(
  evidence = evidence,
  targets  = targets,
  n_sim    = 1000,
  threshold = 0.2
)
```

samp\$results

calc_ciw_2s	<i>Calculates Pre-Posterior Distribution of 95% CI Widths Using Two-step Method</i>
-------------	---

Description

Calculates pre-posterior distribution of 95% CI widths using two-step method.

Usage

```
calc_ciw_2s(N, parms)
```

Arguments

N	A vector of sample sizes
parms	Parameters for the distribution containing: cal_int: calibration intercept cal_slp: calibration slope prev: prevalence dist_type: distribution type estat: c-statistic dist_type: one of ("logitnorm", "beta", "probitnorm") dist_parm1: first parameter of the distribution dist_parm2: second parameter of the distribution

Value

List of length N, of vectors containing 95% confidence interval width for each of: cstat: c-statistic cal_oe: observed to expected ratio cal_mean: mean calibration cal_int: calibration intercept cal_slp: calibration slope

calc_ciw_mc	<i>#'Calculates Pre-Posterior Distribution of 95% CI Widths Based on Given Method</i>
-------------	---

Description

Calculates pre-posterior distribution of 95% CI widths based on given method

Usage

```
calc_ciw_mc(N, parms_sample, method)
```

Arguments

N	A vector of sample sizes
parms_sample	Matrix of parameters for the distribution each row with appropriate parameters: cstat: c-statistic prev: prevalence dist_type: distribution type dist_parm1: first parameter of distribution dist_parm2: second parameter of distribution cal_int: calibration intercept cal_slp: calibration slope
method	Method to calculate 95% confident interval width, one of sample, 2s

Value

List of matrices each with dimension (number of rows in parms_sample x length N) containing 95% confidence interval width for each of: cstat: c-statistic cal_oe: observed to expected ratio cal_mean: mean calibration cal_int: calibration intercept cal_slp: calibration slope

calc_ciw_sample	<i>Calculates Pre-Posterior Distribution of 95% CI Widths Using Sampling-based Simulation</i>
-----------------	---

Description

Calculates pre-posterior distribution of 95% CI widths using sampling-based simulation

Usage

```
calc_ciw_sample(N, parms)
```

Arguments

N	A vector of sample sizes
parms	Parameters for the distribution containing: prev: prevalence dist_type: distribution type dist_parm1: first parameter of distribution dist_parm2: second parameter of distribution cal_int: calibration intercept cal_slp: calibration slope

Value

List of length N, of vectors containing 95% confidence interval width for each of: cstat: c-statistic cal_oe: observed to expected ratio cal_mean: mean calibration cal_int: calibration intercept cal_slp: calibration slope

calc_cstat	<i>Calculates the C-statistic of Model</i>
------------	--

Description

Calculates the c-statistic given the model type and parameters.

Usage

```
calc_cstat(type, parms, m = NULL)
```

Arguments

type	A character string; one of c("beta", "logitnorm", "probitnorm") indicating the model type.
parms	A numeric vector containing parameters relevant to the model.
m	Mean, default is NULL

Value

The C-statistic

calc_riley_vars	<i>Calculates Approximate Variances and Covariance for Performance Metrics</i>
-----------------	--

Description

Calculates approximate variances performance metrics and covariance of calibration intercept and slope using the Riley framework

Usage

```
calc_riley_vars(N, parms)
```

Arguments

N	sample size of the validation dataset
parms	list containing model and distribution parameters: prev: expected prevalence cstat: c-statistic of the model dist_type: one of ("logitnorm", "beta", "probitnorm") dist_parm1: first parameter of the distribution dist_parm2: second parameter of the distribution cal_int: calibration intercept cal_slp: calibration slope

Value

list of approximate variances and covariance of the performance metrics.

calc_se_sp	<i>Calculates the Sensitivity and Specificity</i>
------------	---

Description

Calculate the sensitivity and specificity of the model at given threshold

Usage

```
calc_se_sp(dist_type, dist_parms, cal_int, cal_slp, threshold, prev)
```

Arguments

dist_type	The distribution type, one of c("logitnorm", "beta", "probitnorm").
dist_parms	Vector of the two parameters of interest given the distribution.
cal_int	The calibration intercept.
cal_slp	The calibration slope.
threshold	The risk threshold
prev	The outcome prevalence, the expectation of the model

Value

A vector containing sensitivity and specificity

Examples

```
calc_se_sp("beta", c(1,1), 0.9, 0.75, 0.5, 0.5)
```

find_n_mean	<i>Calculates Sample Size Given Target Mean CI</i>
-------------	--

Description

Calculates sample size N, so that the mean confidence interval is equal to given target, assumes function is decreasing and convex

Usage

```
find_n_mean(target, N, ciws, decreasing = T, convex = T)
```

Arguments

target	The target mean confidence interval width
N	Sample sizes corresponding to each row of ciws,=
ciws	Matrix of confidence intervals widths, each row corresponding to N
decreasing	Logical. Constraining function to decreasing
convex	Logical. Constraining function to convex

Value

Integer. Estimated sample size needed to achieve the target

<code>find_n_quantile</code>	<i>Calculates Sample Size Given Target Quantile</i>
------------------------------	---

Description

Find sample size N, so that the specified quantile is equal to given target

Usage

```
find_n_quantile(target, N, q, ciws)
```

Arguments

target	The desired quantile target value
N	Sample sizes corresponding to each row of ciws
q	Desired quantile level, between 0 and 1.
ciws	A matrix of confidence intervals widths, each row corresponding to N

Value

Estimated sample size needed to achieve the target

infer_cal_int_from_mean

Infer Calibration Intercept from Mean Calibration

Description

Infer calibration intercept from mean calibration given a fixed calibration slope and a given distribution for calibrated risks

Usage

```
infer_cal_int_from_mean(dist_type, dist_parms, cal_mean, cal_slp, prev = NULL)
```

Arguments

dist_type	The distribution type, one of c("logitnorm", "probitnorm", "beta").
dist_parms	The two parameters that index the type.
cal_mean	The mean calibration.
cal_slp	The calibration slope.
prev	Outcome prevalence. Optional; if not provided, estimate is as the expected value of the distribution of calibrated risks.

Value

The estimated calibration intercept

infer_cal_int_from_oe *Infer Calibration Intercept from O/E ratio*

Description

Infer calibration intercept from observed-to-expected outcome ratio given a fixed calibration slope and a given distribution for calibrated risks

Usage

```
infer_cal_int_from_oe(dist_type, dist_parms, cal_oe, cal_slp, prev = NULL)
```

Arguments

dist_type	The distribution type, one of c("logitnorm", "probitnorm", "beta").
dist_parms	The two parameters that index the type.
cal_oe	The observed-to-expected outcome ratio.
cal_slp	The calibration slope.
prev	Outcome prevalence. Optional; if not provided, estimate is as the expected value of the distribution of calibrated risks.

Value

The estimated calibration intercept

infer_correlation	<i>Calculates Correlation</i>
-------------------	-------------------------------

Description

Calculates correlation based on simulated data

Usage

```
infer_correlation(dist_type, dist_parms, cal_int, cal_slp, n, n_sim)
```

Arguments

dist_type	The distribution type
dist_parms	The two parameters of interest for the given distribution type
cal_int	The calibration intercept.
cal_slp	The calibration slope.
n	number of observations for each simulation.
n_sim	number of simulations

Value

correlation among the simulated data

inv_mean_quantile	<i>Calculates the Model Parameters Given Quantile</i>
-------------------	---

Description

Calculate the model parameters given the distribution type, mean, quantile, and percentile.

Usage

```
inv_mean_quantile(type, m, q, p)
```

Arguments

type	The distribution type, one of c("norm", "beta", "logitnorm", "probitnorm").
m	Mean of the of distribution.
q	The quantile value.
p	The percentile at which the quantile occurs.

Value

The model parameters of the given type.

inv_moments	<i>Calculates the Model Parameters Given Moments</i>
-------------	--

Description

Calculates the model parameters of interest given the first two moments.

Usage

```
inv_moments(type, moments)
```

Arguments

type	The distribution type, one of c("norm", "beta", "logitnorm").
moments	A numeric vector containing the first two moments of the model

Value

Returns the two parameters for each model. mean and sd for norm mu and sigma for logitnorm shape1 (alpha) and shape2 (beta) for beta

isaric	<i>Isaric Dataset</i>
--------	-----------------------

Description

Data from the International Severe Acute Respiratory and Emerging Infection Consortium regarding Regions in the UK.

Usage

```
isaric
```

Format

A data frame with 8 rows and 10 columns

Region Region where the sample was drawn

Sample_Size Raw number of total subjects available in the region's dataset

n Number of subjects used in analysis after exclusions

n_events Number of positive subjects

cstat C-statistic

cstat_l Lower bound for the confidence interval of the C-statistic

cal_mean Calibration Mean

cal_mean_l Lower bound for the confidence interval of the calibration mean

cal_slope Calibration slope

cal_slope_l Lower bound of the confidence interval of the calibration slope

Source

Simulated Data

moments

Mean and Variance Calculator

Description

Calculates the first two moments (mean and variance) of the given model type and parameters.

Usage

```
moments(type, parms)
```

Arguments

type The distribution type, one of c("norm", "beta", "logitnorm", "probitnorm").

parms A numeric vector containing parameters relevant to the model.

Value

A numeric vector representing the mean and variance.

plot_cal_distance

Plots Calibration Distance from Simulation Curves

Description

simulates calibration curves based on given method, and uses plot to visualize calibration distance (difference between predicted and observed)

Usage

```
plot_cal_distance(N, sample, method = "loess", X = (1:99)/100)
```

Arguments

N	Number of observations to simulate in each sample
sample	Data frame with columns: dist_type: distribution type dist_parm1: first distribution parameter (e.g. mean, alpha, shape1) dist_parm2: second distribution parameter (e.g. sd, beta, shape2) cal_int: calibration intercept cal_slp: calibration slope
method	One of loess or line, on default is loess
X	Vector of predicted probabilities, on default is 0.01 to 0.99

Value

Plot of simulated calibration curves

Examples

```
sample <- data.frame(
  dist_type = rep("beta", 3),
  dist_parm1 = c(1,2,3),
  dist_parm2 = c(3,4,5),
  cal_int = c(0, 0.05, 0.1),
  cal_slp = c(1, 0.9, 0.8))
plot_cal_distance(N=200, sample=sample)
```

plot_cal_instability *Plots Calibration Instability from Simulated Calibration Curves*

Description

Simulates calibration curves based on given method, and uses plot to visualize calibration instability.

Usage

```
plot_cal_instability(N, sample, method = "loess", X = (1:99)/100)
```

Arguments

N	Number of observations to simulate in each sample
sample	Data frame with columns: dist_type: distribution type dist_parm1: first distribution parameter (e.g. mean, alpha, shape1) dist_parm2: second distribution parameter (e.g. sd, beta, shape2) cal_int: calibration intercept cal_slp: calibration slope
method	One of loess or line, on default is loess
X	Vector of predicted probabilities, on default is 0.01 to 0.99

Value

Plot of simulated calibration curves

Examples

```
sample <- data.frame(
  dist_type = rep("beta", 3),
  dist_parm1 = c(1,2,3),
  dist_parm2 = c(3,4,5),
  cal_int = c(0, 0.05, 0.1),
  cal_slp = c(1, 0.9, 0.8))
plot_cal_instability(N=200, sample=sample)
```

process_evidence	<i>Transforms Evidence Into Standardized Format</i>
------------------	---

Description

Verifies that an evidence object has the required members and standardizes it into a bpm_evidence object. Each element's distribution is recorded together with both its native parameters (\$parms) and its first two moments (\$moments, named m and v).

Usage

```
process_evidence(evidence)
```

Arguments

evidence A named list of evidence elements. The required members are:

- prev: outcome prevalence (defaults to a beta distribution),
- cstat: c-statistic (defaults to a beta distribution),
- cal_slp: calibration slope (defaults to norm), and
- exactly one of cal_mean (mean calibration), cal_oe (observed-to-expected ratio), or cal_int (calibration intercept), each defaulting to norm.

Each element may be given either as a formula, name ~ dist(par1, par2), or as a named list, name = list(type = "dist", ...). The supported distributions (type) are "norm", "beta", "logitnorm", and "probitnorm".

Details

The two parameters of each element may be characterized flexibly, as either native distribution parameters or summary moments. The parameters must be **either all unnamed or all named** (a mix such as beta(0.4, var = 0.04) is ambiguous and raises an error).

Unnamed (positional) parameters are taken as the native parameters of the distribution:

- norm(mean, sd),

- `beta(shape1, shape2)`,
- `logitnorm(mu, sigma)`,
- `probitnorm(mu, sigma)`.

Named parameters are matched against the following aliases (pick one pair per element):

- moments with a variance: mean/var (or m/v),
- moments with a standard deviation: mean/sd (or m/sd),
- a mean and an upper 97.5\
- native beta parameters: alpha/beta,
- native `logitnorm/probitnorm` parameters: mu/sigma.

When moments are supplied, the native parameters are obtained by the method of moments (or, for `cih`, by matching the requested quantile).

Value

A `bpm_evidence` object: the standardized, restructured evidence list.

Examples

```
# Formula form, mixing native parameters and moments:
evidence <- list(
  prev ~ beta(116, 155),      # native beta parameters
  cstat ~ beta(mean = 0.76, sd = 0.006),
  cal_mean ~ norm(-0.009, 0.125),
  cal_slp ~ norm(0.995, 0.024)
)
process_evidence(evidence = evidence)

# Equivalent named-list form:
evidence <- list(
  prev=list(type="beta", mean=0.38, sd=0.2),
  cstat=list(mean=0.7, sd=0.05),
  cal_int=list(mean=0.2, sd=0.2),
  cal_slp=list(mean=0.8, sd=0.3)
)
process_evidence(evidence=evidence)
```

rbnorm

Generates Samples From Normal Distribution

Description

generates samples from a normal distribution using marginal means, variances, and covariance

Usage

```
rbnorm(n, mu1, mu2, var1, var2, cov)
```

Arguments

n	Number of samples to be generated
mu1	Mean of first variable
mu2	Mean of second variable
var1	Variance of first variable
var2	Variance of second variable
cov	Covariance between the two variables

Value

Matrix of nx2 where column 1 contains samples for the first variable, and column 2 contains samples for the second variable conditioned on the first

riley_samp	<i>Calculates Sample Size that Achieves Target CI Widths</i>
------------	--

Description

Calculates sample size that achieves target confidence interval widths using Riley's framework

Usage

```
riley_samp(target_ciws, parms)
```

Arguments

target_ciws	Named list containing target confidence interval width for at least one of: prev: prevalence cstat: c-statistic cal_mean: mean calibration cal_oe: observed to expected outcome ratio cal_int: calibration intercept cal_slp: calibration slope
parms	List containing model parameters and distribution: prev: expected prevalence cstat: c-statistic of the model dist_type: one of ("logitnorm", "beta", "probitnorm") dist_parm1: first parameter of the distribution dist_parm2: second parameter of the distribution cal_int: calibration intercept cal_slp: calibration slope

Value

A named list of estimated sample sizes that achieve target confidence interval widths: fciw.prev, fciw.cstat, fciw.cal_mean, fciw.cal_oe, fciw.cal_int, fciw.cal_slp

Index

* datasets

isaric, 13

bpm_valprec, 2

bpm_valsamp, 4

calc_ciw_2s, 6

calc_ciw_mc, 6

calc_ciw_sample, 7

calc_cstat, 8

calc_riley_vars, 8

calc_se_sp, 9

find_n_mean, 9

find_n_quantile, 10

infer_cal_int_from_mean, 11

infer_cal_int_from_oe, 11

infer_correlation, 12

inv_mean_quantile, 12

inv_moments, 13

isaric, 13

moments, 14

plot_cal_distance, 14

plot_cal_instability, 15

process_evidence, 16

rbnorm, 17

riley_samp, 18