

Package ‘ErrorTracer’

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Type Package

Title Bayesian Error Propagation and Forecast Uncertainty
Decomposition

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Description Provides a full pipeline from regularized or standard regression models (elastic net, linear models, generalized linear models, random forests) to informed Bayesian priors, structured forecast uncertainty decomposition (parameter / environmental / residual, plus a temporal component when the model carries an autocorrelation term), and forecast shelf life analysis (the quantification of when a forecast becomes uninformative). Designed for ecological and genomic forecasting with climate or environmental covariates. Methods build on Bürkner (2017) <doi:10.18637/jss.v080.i01> for Bayesian regression via 'Stan', Friedman, Hastie, and Tibshirani (2010) <doi:10.18637/jss.v033.i01> for elastic net regularization, Wright and Ziegler (2017) <doi:10.18637/jss.v077.i01> for random forests, and Vehtari, Gelman, and Gabry (2017) <doi:10.1007/s11222-016-9696-4> for leave-one-out cross-validation.

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decompose_uncertainty *Extract or recompute uncertainty decomposition*

Description

Returns a data.frame with the uncertainty decomposition stored inside an et_prediction object:

param_var Variance of the posterior linear predictor — captures uncertainty in fitted regression coefficients.

env_var Additional variance arising from measurement or prediction uncertainty in the predictor values (estimated via perturbation in `et_predict`). Zero when `env_noise = NULL`.

residual_var Posterior mean of σ^2 (or its family-specific analogue) — biological process noise, unmeasured drivers, and drift. For autocorrelation models this is the *innovation* variance, not the stationary marginal variance; the autocorrelated accumulation is reported separately in `temporal_var`.

temporal_var (Only present when the model formula contains an autocorrelation term such as `ar()`, `ma()`, `arma()`, `cosy()`, `unstr()`, `sar()`, or `car()`.) Variance attributable to residual temporal or spatial dependence beyond the iid param + residual sum, computed as $\text{pmax}(0, \text{total_var} - (\text{param_var} + \text{residual_var}))$. `env_var` is deliberately excluded from this gap because it is an additive perturbation-based augmentation measured outside of `posterior_predict`.

total_var Variance of the full posterior predictive draws, including any autocorrelation structure modelled by brms.

Usage

```
decompose_uncertainty(predictions, ...)
```

Arguments

<code>predictions</code>	An <code>et_prediction</code> object from <code>et_predict</code> , or an <code>et_prediction_list</code> (grouped).
<code>...</code>	Unused.

Details

All variance components are guaranteed non-negative. When `temporal_var` is present, `param_var + residual_var + temporal_var` reconstructs `total_var` (modulo Monte Carlo error); when it is absent, `param_var + residual_var` does. `env_var` is always additive on top, representing the extra variance that would be contributed by perturbing predictors with `env_noise`.

Value

A `data.frame` with columns `obs_id`, `param_var`, `env_var`, `residual_var`, `total_var` (plus `temporal_var` for autocorrelation models, and a leading group column for grouped predictions).

Examples

```
set.seed(1)
df <- data.frame(y = rnorm(20), x1 = rnorm(20))
fit <- et_fit(y ~ x1, data = df,
             chains = 1, iter = 500, warmup = 250,
             cores = 1, refresh = 0)
new_df <- data.frame(x1 = rnorm(5))
pred <- et_predict(fit, newdata = new_df,
                  env_noise = list(x1 = 0.2),
                  n_draws = 200, n_perturb = 50)
decomp <- decompose_uncertainty(pred)
head(decomp)
```

 et_calibrate

Assess calibration of posterior predictive intervals

Description

Computes observed coverage probability at multiple nominal CI levels. A well-calibrated Bayesian model should produce 90% CIs that contain the true value 90% of the time, etc.

Usage

```
et_calibrate(predictions, observed, response_col = NULL, ci_levels = NULL, ...)
```

Arguments

predictions	An <code>et_prediction</code> or <code>et_prediction_list</code> .
observed	A <code>data.frame</code> with true response values. Must have the same number of rows as <code>predictions\$newdata</code> (rows are matched positionally) and a column with the true response values.
response_col	Character. Name of the response column in <code>observed</code> . Defaults to the left-hand side of the model formula if it can be inferred, otherwise must be specified.
ci_levels	Numeric vector. CI levels to assess. Defaults to all levels present in the <code>et_prediction</code> object.
...	Unused.

Value

A `data.frame` with columns:

ci_level Nominal CI level.

nominal Same as `ci_level`.

observed_coverage Fraction of true values falling inside the CI.

n_obs Number of observations used.

calibration_error Signed difference: `observed - nominal`. Positive = over-coverage (CIs too wide / conservative). Negative = under-coverage (CIs too narrow / overconfident).

sharpness Mean CI width across observations. Sharpness and calibration are complementary: a model can be calibrated but useless if sharpness is poor (very wide CIs).

For grouped predictions, a group column is prepended.

Examples

```

set.seed(1)
df <- data.frame(y = rnorm(20), x1 = rnorm(20))
fit <- et_fit(y ~ x1, data = df,
             chains = 1, iter = 500, warmup = 250,
             cores = 1, refresh = 0)
valid_df <- data.frame(y = rnorm(5), x1 = rnorm(5))
pred <- et_predict(fit, newdata = valid_df,
                  n_draws = 200, n_perturb = 50)
cal <- et_calibrate(pred, observed = valid_df, response_col = "y")
print(cal)

```

et_diagnose

Diagnose a fitted ErrorTracer model

Description

Computes Rhat, effective sample size ratios, divergent transitions, and leave-one-out cross-validation (LOO-CV) for a fitted `et_model`.

Usage

```
et_diagnose(model, loo = TRUE, ...)
```

Arguments

<code>model</code>	An <code>et_model</code> or <code>et_model_list</code> .
<code>loo</code>	Logical. Whether to run LOO-CV (can be slow; default TRUE).
<code>...</code>	Unused.

Value

A list with elements:

convergence List: `rhat_max`, `rhat_all_ok`, `neff_min`, `neff_all_ok`, `n_divergences`.

loo List or NULL: `elpd_loo`, `p_loo`, `looic`, `n_bad_pareto_k`, `loo_object`.

summary Printed summary from `brms::summary()`.

For `et_model_list`, a named list of per-group diagnostic lists plus an aggregated summary `data.frame`.

et_fit

*Fit a Bayesian regression model with informed priors***Description**

Wraps `brms::brm()` and attaches the prior specification, training data reference, and configuration for downstream uncertainty decomposition. Pass priors from `extract_priors` to use regularized-model coefficients as prior means; omit it for default (weakly informative) priors.

Usage

```
et_fit(
  formula,
  data,
  priors = NULL,
  chains = 4L,
  iter = 2000L,
  warmup = floor(iter/2),
  cores = min(chains, parallel::detectCores()),
  seed = 42L,
  adapt_delta = 0.95,
  max_treedepth = 12L,
  grouping = NULL,
  eiv = NULL,
  silent = 2L,
  ...
)
```

Arguments

<code>formula</code>	An R formula, e.g. <code>response ~ .</code> or <code>y ~ x1 + x2</code> .
<code>data</code>	A data frame with all predictors and the response.
<code>priors</code>	An <code>et_prior_spec</code> object from <code>extract_priors</code> , or a <code>brmsprior</code> object, or <code>NULL</code> for <code>brms</code> defaults.
<code>chains</code>	Integer. Number of MCMC chains (default 4).
<code>iter</code>	Integer. Total iterations per chain, including warmup (default 2000).
<code>warmup</code>	Integer. Warmup iterations per chain (default <code>floor(iter / 2)</code>).
<code>cores</code>	Integer. Parallel cores (default <code>min(chains, parallel::detectCores())</code>).
<code>seed</code>	Integer. Random seed for reproducibility (default 42).
<code>adapt_delta</code>	Numeric. Target acceptance probability for HMC (default 0.95).
<code>max_treedepth</code>	Integer. Maximum tree depth (default 12).
<code>grouping</code>	Character. Name of a column in <code>data</code> to use for grouping. If non- <code>NULL</code> , one model is fitted per unique group value and an <code>et_model_list</code> is returned.

eiv	Optional errors-in-variables specification. A named list/ vector mapping predictor names to either a scalar SD or a vector of per-row SDs (length nrow(data)). For each entry, the formula term for that predictor is rewritten as <code>brms::me(pred, se_pred)</code> (an auxiliary <code>se_<pred></code> column is appended to data), so the posterior reflects measurement error in the predictor as well as coefficient uncertainty. The beta posteriors widen accordingly, which partially absorbs what ErrorTracer's downstream <code>env_var</code> component would otherwise report. When <code>eiv</code> is supplied together with an <code>et_prior_spec</code> from <code>extract_priors</code> , the informed priors are <i>dropped</i> because they target <code>class = "b"</code> terms and <code>me()</code> terms live under <code>class = "bsp"</code> ; <code>brms</code> defaults are used instead (and a warning is logged).
silent	Integer passed to <code>brms::brm()</code> (default 2, no Stan output).
...	Additional arguments passed to <code>brms::brm()</code> .

Value

An `et_model` object (or an `et_model_list` if grouping is specified).

Examples

```
set.seed(1)
df <- data.frame(y = rnorm(20), x1 = rnorm(20), x2 = rnorm(20))
ps <- extract_priors(lm(y ~ x1 + x2, data = df))
fit <- et_fit(y ~ x1 + x2, data = df, priors = ps,
             chains = 1, iter = 500, warmup = 250,
             cores = 1, refresh = 0)
print(fit)
```

`et_plot_calibration` *Plot calibration: observed vs nominal coverage*

Description

A well-calibrated model produces points along the 1:1 diagonal. Points above the diagonal indicate over-coverage (conservative); below indicates under-coverage (anti-conservative).

Usage

```
et_plot_calibration(cal, group_col = NULL)
```

Arguments

cal	A data.frame from <code>et_calibrate</code> .
group_col	Optional character. Name of a column in <code>cal</code> that identifies sub-groups (e.g. "species", "cluster_id"). When NULL (default), <code>et_plot_calibration</code> uses the group column if present; otherwise it auto-detects any single non-canonical column with more than one unique value and treats it as the grouping. Set to NA to force a single un-grouped series.

Value

A ggplot2 object.

et_plot_coefficients *Forest plot of regression coefficients*

Description

Compares Bayesian posterior estimates (95% CI) with the regularized coefficient values used as prior means (shown as crosses).

Usage

```
et_plot_coefficients(model)
```

Arguments

model An et_model or et_model_list.

Value

A ggplot2 object.

et_plot_decomposition *Plot uncertainty decomposition*

Description

Produces a stacked bar chart showing the relative contributions of parameter, environmental, and residual variance for each observation, plus a fourth temporal-autocorrelation component when present in decomp (see [decompose_uncertainty](#)).

Usage

```
et_plot_decomposition(decomp, proportional = TRUE, group_col = NULL)
```

Arguments

decomp A data.frame from [decompose_uncertainty](#) or directly the \$decomposition slot of an et_prediction.

proportional Logical. If TRUE (default), bars are scaled to sum to 1 (proportional contribution). If FALSE, raw variances are shown.

group_col Character. Optional name of a grouping column in decomp (present for grouped predictions).

Value

A ggplot2 object.

et_plot_forecast	<i>Plot posterior predictive fan chart</i>
------------------	--

Description

Overlays nested credible interval ribbons on the median forecast, with optional observed values for calibration assessment.

Usage

```
et_plot_forecast(  
  predictions,  
  observed = NULL,  
  response_col = NULL,  
  time_col = NULL  
)
```

Arguments

predictions	An et_prediction object.
observed	Optional data.frame with true response values. If provided, points are overlaid.
response_col	Character. Name of the response column in observed.
time_col	Character. Column in predictions\$newdata used as the x-axis. Defaults to observation index.

Value

A ggplot2 object.

et_plot_prior_posterior	<i>Plot prior vs posterior distributions for model coefficients</i>
-------------------------	---

Description

Overlays prior and posterior density for each predictor coefficient, visualising how much the data update the priors.

Usage

```
et_plot_prior_posterior(model, max_preds = 8L, n_prior_draws = 4000L)
```

Arguments

model	An et_model object.
max_preds	Integer. Maximum number of predictors to show (default 8). Predictors are shown in the order they appear in the prior specification.
n_prior_draws	Integer. Number of random draws for the prior density (default 4000).

Value

A ggplot2 object.

et_plot_sensitivity *Plot a sensitivity profile*

Description

Visualises the output of [et_sensitivity_profile](#): for each noise grid point, shows how the **environmental share** of total variance and the **forecast horizon** respond. Observed horizons are drawn as solid points; projected horizons are hollow points; lower-bound rows are drawn as upward arrows at the last informative time.

Usage

```
et_plot_sensitivity(sens, show = c("horizon", "env_share", "ratio"))
```

Arguments

sens	An et_sensitivity object from et_sensitivity_profile .
show	"horizon" (default) shows the shelf-life horizon; "env_share" shows env_var / total_var; "ratio" shows the mean CI width / plausible range.

Details

The x-axis is the noise fraction (when the grid was built from fraction_grid) or the grid step label otherwise. When both a numeric fraction and a descriptive label exist, the fraction is preferred for continuous x-positioning.

Value

A ggplot2 object.

See Also

[et_sensitivity_profile](#)

et_plot_shelf_life	<i>Plot forecast shelf life</i>
--------------------	---------------------------------

Description

Shows how credible interval width grows over the forecast horizon and marks the threshold beyond which the forecast is uninformative.

Usage

```
et_plot_shelf_life(sl, show_ratio = TRUE)
```

Arguments

sl	An <code>et_shelf_life</code> object from shelf_life .
show_ratio	Logical. If TRUE (default), plots the ratio (CI width / plausible range) rather than raw CI width.

Value

A `ggplot2` object.

et_predict	<i>Posterior prediction with uncertainty decomposition</i>
------------	--

Description

Generates posterior predictive draws for new observations, propagates environmental measurement uncertainty through the model, and computes credible intervals. The resulting `et_prediction` object is the input to [decompose_uncertainty](#), [shelf_life](#), and the plotting functions.

Usage

```
et_predict(
  model,
  newdata,
  env_noise = NULL,
  env_cov = NULL,
  env_dist = NULL,
  n_draws = 2000L,
  ci_levels = c(0.5, 0.8, 0.9, 0.95),
  n_perturb = NULL,
  n_env_draws = 1L,
  interval_type = c("predictive", "linpred"),
  include_env_in_ci = FALSE,
  ...
)
```

Arguments

model	An <code>et_model</code> or <code>et_model_list</code> object from <code>et_fit</code> .
newdata	A <code>data.frame</code> containing the predictor columns named in the model formula. For grouped models, must also contain the grouping column.
env_noise	Environmental measurement / prediction uncertainty. Can be: <ul style="list-style-type: none"> • NULL (default): no environmental noise. • A single numeric: applied as a fraction of each predictor's empirical SD in <code>newdata</code> (e.g. <code>0.1</code> means 10% noise, constant across all observations). • A named list or named numeric vector with one scalar per predictor: constant absolute noise SD per predictor, e.g. <code>list(Tmean = 0.5, PPT = 10)</code>. • A named list where each entry is a numeric vector of length <code>nrow(newdata)</code>: <i>time-varying</i> (per-row) noise SDs. Use this when predictor uncertainty grows with forecast horizon, e.g. from a GCM ensemble spread that increases over time: <code>list(Tmean = 0.30 + 0.01 * (years - base_year))</code>. Entries not supplied default to zero (no noise for that predictor).
env_cov	Correlation structure of the environmental noise. The <i>magnitudes</i> of the noise come from <code>env_noise</code> ; <code>env_cov</code> supplies the <i>correlation</i> between predictors, so that a perturbation on row i is drawn from $\mathcal{N}(0, D_i R D_i)$ with $D_i = \text{diag}(\sigma_{i1}, \dots, \sigma_{ip})$ and R = the correlation matrix. One of: <ul style="list-style-type: none"> • NULL (default): independent noise, $R = I$ — equivalent to ErrorTracer behaviour prior to this feature and the right choice when predictor measurement errors are genuinely independent (e.g. separate instruments on unrelated variables). • "empirical": compute the correlation of the predictor columns in the training data (<code>model\$data</code>). Use this when predictor <i>errors</i> are expected to inherit the correlation structure of the predictors themselves — e.g. temperature and humidity that co-vary in the underlying climate system. • "newdata": compute the correlation of the predictor columns in <code>newdata</code>. Useful when the forecast window has a different covariance structure than training (e.g. scenario runs). • A numeric $p \times p$ matrix with <code>dimnames</code> matching the model's predictors. Entries with an off-diagonal exceeding 1 are rescaled to a correlation matrix. Use this to supply an independent estimate of the <i>error</i> correlation structure (e.g. from a reanalysis product or a sensor covariance report). <p>A correlation derived from training data is a working assumption: the structure of the <i>errors</i> is assumed to mirror the structure of the <i>values</i>. When this is implausible, pass a matrix directly.</p>
env_dist	Distributional form of the per-predictor noise. The <code>env_noise</code> SDs set the <i>magnitude</i> of the perturbation; <code>env_dist</code> sets its <i>shape</i> . For every distribution other than "gaussian", the noise is calibrated so that (approximately) $E[\hat{x}] = x$ and $\text{Var}[\hat{x}] = \sigma^2$, using a Gaussian copula to honour <code>env_cov</code> . One of: <ul style="list-style-type: none"> • NULL (default): "gaussian" for every predictor — additive Gaussian noise, legacy behaviour. • A single string ("gaussian", "lognormal", "gamma", "beta"): applied to all predictors.

- A named list / character vector with one entry per predictor to override the default, e.g. `list(PPT = "gamma", tmax = "gaussian")`.

Distributions:

"gaussian" Additive normal noise ($\tilde{x} = x + \varepsilon, \varepsilon \sim N(0, \sigma^2)$). Appropriate for symmetric measurement error on a continuous, potentially negative scale (temperature, anomalies).

"lognormal" Multiplicative noise: $\log \tilde{x} \sim N(\log x - s^2/2, s^2)$ with $s^2 = \log(1 + (\sigma/x)^2)$. Preserves positivity; right-tail skewed. Natural for strictly positive continuous variables whose error scales with magnitude (e.g. enzyme activity, biomass). Rows with $x \leq 0$ are left unperturbed.

"gamma" $\tilde{x} \sim \text{Gamma}(\text{shape} = (x/\sigma)^2, \text{rate} = x/\sigma^2)$. Positive support, right-skewed, analytic mean/variance match. Natural for precipitation, rates, and other non-negative continuous variables. Rows with $x \leq 0$ are left unperturbed.

"beta" $\tilde{x} \sim \text{Beta}(\alpha, \beta)$ with $\alpha + \beta = x(1 - x)/\sigma^2 - 1$. Support in $(0, 1)$; appropriate for proportions and probabilities (allele frequencies, presence rates). Rows with $x \notin (0, 1)$ or $\sigma^2 \geq x(1 - x)$ are left unperturbed.

Correlation (`env_cov`) is applied to the latent standard-normal draws before the marginal quantile transform, so rank correlations are preserved across distributions.

<code>n_draws</code>	Integer. Number of posterior draws to use (default 2000; capped at the number of draws available in the fit).
<code>ci_levels</code>	Numeric vector. Credible interval levels to compute (default <code>c(0.5, 0.8, 0.9, 0.95)</code>).
<code>n_perturb</code>	Integer. Number of posterior draws used for the environmental perturbation step (default <code>min(500, n_draws)</code>). Reducing this speeds up computation.
<code>n_env_draws</code>	Integer. Number of independent environmental perturbations averaged <i>per posterior draw</i> when estimating <code>env_var</code> (default 1). Increasing this reduces Monte Carlo noise on the environmental-variance estimate at the cost of proportional computation. The decomposition always reports a Monte Carlo SE (<code>v_env_mcse</code>) alongside <code>env_var</code> ; it decreases roughly like $1/\sqrt{n_env_draws \cdot n_perturb}$.
<code>interval_type</code>	Character. Which draws to use when computing credible intervals: <ul style="list-style-type: none"> • "predictive" (default): draws from <code>posterior_predict</code>, which include sigma (residual noise). Use this when forecasting individual observations — e.g. a single population's allele frequency, one site's ozone reading on a specific day. • "linpred": draws from <code>posterior_linpred</code>, which capture only parameter uncertainty (no sigma). Use this when forecasting the mean response — e.g. the expected ozone across many similar days, or mean delta f across replicate populations. These intervals are always narrower; they will under-cover individual observations unless sigma is negligible. <p>The decomposition components and <code>posterior_predict</code> / <code>posterior_linpred</code> matrices are always computed regardless of this setting.</p>
<code>include_env_in_ci</code>	Logical. When TRUE and <code>interval_type = "predictive"</code> , credible intervals are constructed from environmentally inflated draws $\tilde{y} = \tilde{\mu} + \varepsilon$, with $\tilde{\mu}$ the

perturbed linear predictor and $\varepsilon \sim N(0, \sigma^2)$ using posterior draws of σ . This folds the environmental uncertainty component back into the CI, which is typically what you want for sensitivity analyses or whenever the reported interval should cover predictor-measurement error. When FALSE (default, backward compatible), CIs are based on posterior_predict only — parameter + residual, without predictor noise.

... Passed to methods.

Value

An et_prediction object (list) containing:

posterior_predict Matrix [n_draws x n_obs]: full posterior predictive draws (parameter + residual uncertainty).

posterior_linpred Matrix [n_draws x n_obs]: linear predictor draws on the **link scale** (parameter uncertainty only).

lp_perturbed Matrix [n_perturb x n_obs]: linear predictor on the link scale computed on environmentally perturbed inputs.

sigma_draws Numeric vector: posterior draws of sigma (NA for families without a sigma parameter, e.g. \ Binomial).

credible_intervals data.frame with columns row_id, ci_level, lower, median, upper, width.

decomposition data.frame with columns obs_id, total_var, param_var, env_var, v_env_mcse, residual_var. All components are on the **response scale**. v_env_mcse is the Monte Carlo SE of env_var. residual_var is per-observation for non-Gaussian families (e.g. \ Binomial: $E_s[\mu^{(s)}(1 - \mu^{(s)})]$) and constant for Gaussian.

newdata The input newdata.

model Reference to the et_model used.

env_cov The $p \times p$ correlation matrix actually used for perturbation (identity for env_cov = NULL).

env_dist Named character vector mapping each predictor to the distribution actually used for its perturbation.

See Also

[decompose_uncertainty](#), [shelf_life](#), [et_calibrate](#)

et_sensitivity_profile

Sensitivity profile of the environmental uncertainty component

Description

Sweeps a grid of noise magnitudes, re-runs [et_predict](#) for each, and summarises how the environmental variance component and the forecast shelf life respond. This turns the subjective choice of env_noise into an auditable *curve*: instead of reporting a single shelf life at one (often hand-picked) measurement-error level, the user sees how the horizon degrades as assumed noise grows.

Usage

```

et_sensitivity_profile(
  model,
  newdata,
  response_scale,
  fraction_grid = NULL,
  absolute_grid = NULL,
  env_cov = NULL,
  env_dist = NULL,
  include_env_in_ci = TRUE,
  ci_level = 0.9,
  ci_levels = c(0.5, 0.8, 0.9, 0.95),
  threshold = 1,
  time_col = NULL,
  n_draws = 2000L,
  n_perturb = NULL,
  max_extrapolation_factor = 10,
  verbose = TRUE,
  plausible_range = NULL
)

```

Arguments

model	An <code>et_model</code> from <code>et_fit</code> .
newdata	<code>data.frame</code> passed to <code>et_predict</code> .
response_scale	Two-element numeric vector <code>c(min, max)</code> , as in <code>shelf_life</code> . The effective range is <code>diff(response_scale)</code> .
fraction_grid	Optional numeric vector of scalar noise fractions.
absolute_grid	Optional list of <code>env_noise</code> arguments (each a named list / vector). If supplied together with <code>fraction_grid</code> , <code>fraction_grid</code> is ignored.
env_cov, env_dist	Passed through to <code>et_predict</code> .
include_env_in_ci	Logical. If <code>TRUE</code> (default), credible intervals — and hence the shelf-life horizon — are computed from environmentally inflated draws (<code>et_predict(..., include_env_in_ci = TRUE)</code>). Set <code>FALSE</code> to see how <code>env_var</code> evolves while holding the CI fixed at parameter + residual.
ci_level	Numeric. Used for shelf life and CI width tracking (default 0.90). Must be in <code>ci_levels</code> .
ci_levels	Numeric vector of CI levels computed per iteration (default <code>c(0.5, 0.8, 0.9, 0.95)</code>).
threshold	Numeric. Shelf-life threshold (default 1.0).
time_col	Character. Optional time column for shelf life.
n_draws, n_perturb	Passed to <code>et_predict</code> .

max_extrapolation_factor	Passed to shelf_life.
verbose	Logical. If TRUE (default), logs each iteration.
plausible_range	Deprecated. Use response_scale instead.

Details

Three argument styles are supported for the grid:

- `fraction_grid`: scalar fractions of each predictor's SD in newdata. These are passed straight to the scalar form of `et_predict`'s `env_noise`. Use this for an apples- to-apples sweep that scales all predictors together.
- `absolute_grid`: a list of named numeric lists / vectors; each list element becomes a single `env_noise` argument (so you can sweep absolute SDs for one predictor while holding others fixed).
- Neither supplied (NULL): a default log-spaced fraction grid $c(0, 0.01, 0.025, 0.05, 0.1, 0.2, 0.4, 0.8)$ is used.

Each iteration calls `et_predict(..., env_noise = g)` then `shelf_life` and records:

- Mean parameter / environmental / residual / total variance across forecast observations.
- The horizon description ("observed", "projected", or "lower_bound") and numeric horizon if any.
- Mean / max CI width / plausible-range ratio.

Value

An `et_sensitivity` object: a data.frame with one row per grid point and columns

`grid_id` 1-indexed step.

`label` Short descriptor of the `env_noise` used (e.g. "fraction = 0.05").

`fraction` Scalar fraction when applicable; NA for absolute-grid rows.

`param_var`, `env_var`, `residual_var`, `total_var` Mean across forecast observations.

`env_share` $\text{env_var} / (\text{env_var} + \text{total_var})$ — the fraction of combined predictive variance attributable to predictor noise (bounded in $[0, 1]$).

`ci_width_mean`, `ci_width_max` At `ci_level`.

`ratio_mean`, `ratio_max` CI width / plausible range.

`horizon_type` One of "observed", "projected", "lower_bound".

`horizon` Numeric horizon (observed or projected) or NA for lower-bound rows.

`horizon_description` The long description returned by `shelf_life`.

The returned object carries the original call and `response_scale` as attributes for use by `et_plot_sensitivity`.

See Also

[et_predict](#), [shelf_life](#), [et_plot_sensitivity](#)

et_sim

*Simulated allele-frequency-change dataset for ErrorTracer tutorials***Description**

A named list containing training data, forecast predictors, validation responses, and ground-truth parameters for two synthetic SNP clusters (A and B) at a hypothetical mountain plant site. The dataset is designed to exercise every step of the ErrorTracer pipeline and to support parameter-recovery validation (true coefficients are known).

Usage

et_sim

Format

A named list with seven elements:

`train` `data.frame` with 100 rows and 6 columns (2 clusters \times 50 training years, 1970–2019):

year Integer. Calendar year.

cluster_id Character. SNP cluster identifier: "A" or "B".

Tmean Numeric. Standardised mean growing-season temperature (original unit: °C; standardised using training-period mean and SD).

PPT Numeric. Standardised total growing-season precipitation (original unit: mm).

SWE Numeric. Standardised peak snow water equivalent (original unit: mm).

z_diff Numeric. Simulated allele-frequency change on the arcsin-sqrt scale ($\arcsin(\sqrt{f})$ transformation). This transformation is unbounded and has no fixed $[-1, 1]$ constraint; the plausible range should be derived from the training data or from the theoretical bounds $([0, \pi/2] \setminus \{0\})$ on the arcsin scale).

`forecast` `data.frame` with 30 rows and 5 columns (2 clusters \times 15 forecast years, 2020–2034).

Same columns as `train` except `z_diff` is absent — these are the prediction targets. Predictors are standardised using the training-period statistics stored in `standardization`.

`validation` `data.frame` with 30 rows and 3 columns (`year`, `cluster_id`, `z_diff`). True response values for the forecast period; used with `et_calibrate` to assess posterior predictive coverage.

`true_params` Named list with one element per cluster. Each element is a named numeric vector of the true data-generating parameters: `intercept`, `Tmean`, `PPT`, `SWE`, and `sigma` (residual SD).

Cluster A `intercept = 0.00`, `Tmean = 0.50`, `PPT = -0.30`, `SWE = 0.20`, `sigma = 0.30`

Cluster B `intercept = 0.10`, `Tmean = 0.30`, `PPT = -0.20`, `SWE = -0.25`, `sigma = 0.35`

`env_noise` Named list. Suggested per-predictor environmental noise SDs (on the standardised scale) for use with the `env_noise` argument of `et_predict`: `Tmean = 0.30`, `PPT = 0.20`, `SWE = 0.15`.

standardization Named list with one element per predictor. Each element is a named numeric vector `c(mean = ..., sd = ...)` giving the training-period statistics used to standardise the data. Needed if you wish to back-transform predictions to the original (unstandardised) scale with `unstandardize`.

description Character string. Human-readable description of the dataset and how it was generated.

Details

The climate time series are simulated as AR(1) processes with a warming trend in T_{mean} ($+0.015$ °C yr^{-1} , cumulating to ~ 0.30 SD above the training mean over the 15-year forecast window). SWE is generated with a weak dependence on T_{mean} (negative coupling) and PPT (positive coupling) plus a dominant independent noise component, so that the three predictors have only mild pairwise correlation ($|R| < 0.2$ on the training subset). This makes all regression coefficients reliably identifiable in a single replicate. All predictors are standardised using training-period statistics only.

In addition to the three real climate predictors, the dataset includes ten independent standardised nuisance predictors d_1, \dots, d_{10} with zero true effect on the response. They give the regularized-prior extraction step a real selection job: `cv.glmnet` ($\alpha = 0.5$) is expected to identify the three real predictors and shrink the ten dummies toward zero. Without them, regularization on three well-identified predictors merely biases the true coefficients without selecting anything.

Responses are generated from a linear model with Gaussian noise:

$$z_diff_t = \alpha + \beta_1 T_{\text{mean}_t} + \beta_2 PPT_t + \beta_3 SWE_t + \varepsilon_t, \quad \varepsilon_t \sim \mathcal{N}(0, \sigma^2)$$

The true parameters are stored in `et_sim>true_params` for parameter-recovery validation.

The dataset was generated with `set.seed(111)`. The full generation script is at `data-raw/generate_et_sim.R`.

Source

Generated by `data-raw/generate_et_sim.R` with `set.seed(111)`.

Examples

```
data(et_sim)

# Inspect structure
str(et_sim, max.level = 2)

# Training data
head(et_sim$train)

# True parameters for cluster A
et_sim>true_params$A

# Suggested noise SDs for et_predict()
et_sim$env_noise
```

et_theme	<i>Minimal ggplot2 theme for ErrorTracer plots</i>
----------	--

Description

Minimal ggplot2 theme for ErrorTracer plots

Usage

```
et_theme(base_size = 12)
```

Arguments

base_size Base font size (default 12).

Value

A ggplot2 theme object.

extract_priors	<i>Extract brms prior specification from a regularized or standard regression model</i>
----------------	---

Description

Converts a fitted model object into a brms prior specification suitable for `et_fit`. The coefficient estimates (or importance weights for **ranger**) from the regularized or standard fit are used as informative prior means, so the Bayesian model starts close to the regularized or standard solution while remaining open to data-driven revision.

Usage

```
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
  intercept_prior_sd = NULL,
  sigma_prior_scale = 1,
  ...
)

## S3 method for class 'lm'
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
```

```
    intercept_prior_sd = NULL,
    sigma_prior_scale = 1,
    ...
)

## S3 method for class 'glm'
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
  intercept_prior_sd = NULL,
  sigma_prior_scale = 1,
  ...
)

## S3 method for class 'cv.glmnet'
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
  intercept_prior_sd = NULL,
  sigma_prior_scale = 1,
  lambda = "lambda.min",
  ...
)

## S3 method for class 'glmnet'
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
  intercept_prior_sd = NULL,
  sigma_prior_scale = 1,
  s = 1L,
  ...
)

## S3 method for class 'ranger'
extract_priors(
  model,
  multiplier = 2,
  min_sd = 0.1,
  intercept_prior_sd = NULL,
  sigma_prior_scale = 1,
  ...
)
```

Arguments

model	A fitted model. Supported classes: <ul style="list-style-type: none"> • <code>cv.glmnet / glmnet</code> — elastic net / lasso • <code>lm</code> — ordinary least squares • <code>glm</code> — generalized linear model • <code>ranger</code> — random forest (importance-scaled flat priors)
multiplier	Numeric scalar. Prior SD is set to <code>multiplier * coef </code> (for signed-coefficient methods) or <code>multiplier * importance_normalised</code> (for <code>ranger</code>). Default 2.0.
min_sd	Numeric scalar. Minimum prior SD to avoid degenerate (spike) priors on near-zero coefficients. Default 0.1.
intercept_prior_sd	Optional prior SD for the intercept term. The default NULL emits <i>no</i> explicit intercept prior and lets <code>brms</code> pick its data-aware default (<code>student_t(3, median(y), mad(y))</code>). If you supply a numeric value, the prior becomes <code>normal(0, intercept_prior_sd)</code> — this is only appropriate when the response has been centred (or nearly so) before fitting, since <code>brms</code> 's <code>class = "Intercept"</code> refers to the intercept at the centre of the predictors, whose posterior mean equals $E[y]$ there. Supplying a small <code>intercept_prior_sd</code> for an uncentred response will pull the intercept toward zero and cripple the fit.
sigma_prior_scale	Scale parameter for the half-Cauchy prior on the residual SD <code>sigma</code> . Default 1.0.
...	Additional arguments passed to methods.
lambda	Which lambda to extract coefficients from when the model is a <code>cv.glmnet</code> object. Either <code>"lambda.min"</code> (default) or <code>"lambda.1se"</code> .
s	Index (integer) or value of lambda to extract coefficients at when the model is a plain <code>glmnet</code> object. Defaults to the first lambda (smallest regularisation).

Details

For `ranger` models, signed coefficients are not available. Priors are centred at zero (direction unknown) and the prior SD for each predictor is set to `multiplier * importance_normalised`, where `importance` is normalised to the `[min_sd, 1]` interval. Only variables with positive permutation importance are included.

Value

An `et_prior_spec` list containing:

`prior` A `brmsprior` object for use in `brms::brm()`.

`pred_names` Character vector of included predictor names.

`coefs` Named numeric vector of regularized coefficients (NULL for `ranger`, which uses importance instead).

`method` Character: the dispatch method used.

`multiplier, min_sd` Settings echo.

Examples

```
fit_lm <- lm(mpg ~ wt + hp + cyl, data = mtcars)
ps <- extract_priors(fit_lm, multiplier = 2, min_sd = 0.1)
print(ps)
```

shelf_life

*Compute the forecast shelf life***Description**

Quantifies *when* a forecast becomes uninformative by comparing the width of credible intervals to a response scale. A forecast is uninformative when its CI width exceeds `threshold * response_scale`.

Usage

```
shelf_life(
  predictions,
  response_scale,
  ci_level = 0.9,
  threshold = 1,
  time_col = NULL,
  min_slope_for_projection = 1e-04,
  max_extrapolation_factor = 10,
  ...,
  plausible_range = NULL
)
```

Arguments

<code>predictions</code>	An <code>et_prediction</code> or <code>et_prediction_list</code> .
<code>response_scale</code>	Numeric vector of length 2 (<code>c(min, max)</code>) giving the response scale used as the denominator in the CI-width / range ratio. For unbounded responses use <code>range(training_data\$response)</code> as a conservative default, or supply a biologically motivated interval. The effective range is <code>diff(response_scale)</code> .
<code>ci_level</code>	Numeric. The credible interval level to use (default 0.90). Must be present in the <code>et_prediction</code> object.
<code>threshold</code>	Numeric. CI width / response scale above which the forecast is uninformative (default 1.0).
<code>time_col</code>	Character. Optional column in <code>predictions\$newdata</code> to use as the time axis. If NULL, observation index is used.
<code>min_slope_for_projection</code>	Numeric. Minimum linear slope (of ratio vs. time) required to attempt extrapolation when all periods are informative. Below this value the shelf life is reported as a lower bound only. Default 1e-4.

max_extrapolation_factor
 Numeric. Cap on how far the linear projection may reach beyond the observed window. If the projected crossing time exceeds $\max(\text{time}) + \text{max_extrapolation_factor} * (\max(\text{time}) - \min(\text{time}))$, the result is reported as a lower bound instead of a projection. Set to Inf to disable the cap. Default 10.

...
 plausible_range
 Unused.
 Deprecated. Use response_scale instead.

Details

The function operates in three modes depending on the available data and whether the uninformative threshold is crossed within the forecast window:

Observed The threshold is crossed within the forecast/validation window. The shelf life is the first time point at which $\text{ratio} \geq \text{threshold}$.

Projected All forecast periods remain informative but the CI/range ratio is trending upward. A linear trend is fitted to the ratios and extrapolated to estimate when the threshold would be reached. The projected crossing time $t^* = (\tau - a)/b$ (where τ is the threshold, a the fitted intercept, b the fitted slope) is reported together with a Monte Carlo standard error se_t_star derived via the delta method.

Lower bound All forecast periods are informative with no upward trend in the ratio. The shelf life is reported as a lower bound: $>$ last observed time.

The intended workflow is:

1. Fit the model ([et_fit](#)).
2. Predict on held-out data or a future time window ([et_predict](#)).
3. Call `shelf_life()` on those predictions.
4. If the threshold is not crossed in the held-out window, the *projected* mode extrapolates the horizon automatically.

Value

An `et_shelf_life` object (a `data.frame`) with columns:

obs_id Observation index.
time Time axis value.
ci_width Width of the credible interval at `ci_level`.
plausible_range Effective response scale (scalar diff).
ratio CI width / response scale.
informative Logical; TRUE when $\text{ratio} < \text{threshold}$.

For grouped predictions a group column is prepended. The object carries three attributes that drive `print()`:

`horizon` Named list with elements `value`, `type` ("observed", "projected", or "lower_bound"), `last_informative`, `description`, and (for projected) `se_t_star`.
`horizon_by_group` For grouped objects: named list of per-group horizon lists.
`threshold` The threshold value used.

Examples

```

set.seed(1)
df <- data.frame(y = rnorm(20), year = 2001:2020, x1 = rnorm(20))
fit <- et_fit(y ~ x1, data = df,
             chains = 1, iter = 500, warmup = 250,
             cores = 1, refresh = 0)
new_df <- data.frame(x1 = rnorm(10), year = 2021:2030)
pred <- et_predict(fit, newdata = new_df,
                  n_draws = 200, n_perturb = 50)
sl <- shelf_life(pred,
                 response_scale = range(df$y),
                 ci_level = 0.90,
                 threshold = 1.0,
                 time_col = "year",
                 max_extrapolation_factor = 10)

print(sl)

```

standardize	<i>Standardize a numeric vector (mean-centre, unit-variance)</i>
-------------	--

Description

Standardize a numeric vector (mean-centre, unit-variance)

Usage

```
standardize(x)
```

Arguments

x Numeric vector.

Value

Standardized numeric vector. Returns zeros if variance is zero.

unstandardize	<i>Reverse standardization</i>
---------------	--------------------------------

Description

Reverse standardization

Usage

```
unstandardize(z, mu, s)
```

Arguments

<i>z</i>	Standardized values.
<i>mu</i>	Original mean.
<i>s</i>	Original standard deviation.

Value

Values on the original scale.

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