

Package: palimpsestr (via r-universe)

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Type Package

Title Probabilistic Decomposition of Archaeological Palimpsests

Version 0.12.0

Description Probabilistic framework for the analysis of archaeological palimpsests based on the Stratigraphic Entanglement Field (SEF). Integrates spatial proximity, stratigraphic depth, chronological overlap, and cultural similarity to estimate latent depositional phases via diagonal Gaussian mixture Expectation-Maximisation (EM). Provides the Stratigraphic Entanglement Index (SEI), Excavation Stratigraphic Energy (ESE), and Palimpsest Dissolution Index (PDI) for quantifying depositional coherence, detecting intrusive finds, and measuring palimpsest formation. Includes simulation, diagnostics, phase-count selection, publication-quality plots, and Geographic Information System (GIS) export via 'sf'. Methods are described in Cocca (2026)
<<https://github.com/enzococca/palimpsestr>>.

URL <https://github.com/enzococca/palimpsestr>

BugReports <https://github.com/enzococca/palimpsestr/issues>

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`adjusted_rand_index` *Adjusted Rand Index*

Description

Compares estimated phase assignments against known true labels using the Adjusted Rand Index (Hubert and Arabie, 1985). Values near 1 indicate perfect agreement; values near 0 indicate random agreement.

Usage

```
adjusted_rand_index(object, true_labels)
```

Arguments

`object` A `sef_fit` object, or an integer vector of predicted labels.
`true_labels` Integer vector of known phase assignments.

Value

A single numeric value in $[-1, 1]$.

See Also

[confusion_matrix](#), [fit_sef](#)

Other validation: [bootstrap_sef\(\)](#), [confusion_matrix\(\)](#), [cv_sef\(\)](#), [optimize_weights\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1, mixing = 0.05)
fit <- fit_sef(x, k = 3, seed = 1)
adjusted_rand_index(fit, x$true_phase)
```

archaeo_sim

Simulate an archaeological palimpsest dataset

Description

Generates a synthetic excavation dataset with known latent phases, controlled spatial clustering, and configurable inter-phase mixing.

Usage

```
archaeo_sim(n = 150, k = 3, seed = NULL, mixing = 0.08)
```

Arguments

n	Number of observations (finds).
k	Number of latent depositional phases.
seed	Optional random seed for reproducibility.
mixing	Proportion of observations to perturb spatially and taphonomically, simulating post-depositional disturbance (0–1).

Value

A data.frame with columns: id, x, y, z, context, date_min, date_max, class, taf_score, true_phase.

See Also

[fit_sef](#) for fitting the SEF model to the output.

Examples

```
easy <- archaeo_sim(n = 100, k = 3, mixing = 0.05, seed = 1)
table(easy$true_phase)

hard <- archaeo_sim(n = 200, k = 4, mixing = 0.50, seed = 1)
table(hard$true_phase)
```

as_phase_table	<i>Extract phase probability table</i>
----------------	--

Description

Returns a data.frame combining dominant phase assignments, membership probabilities, entropy, local SEI, and energy.

Usage

```
as_phase_table(object)
```

Arguments

object A sef_fit object.

Value

A data.frame with one row per find.

See Also

[predict_phase](#), [phase_diagnostic_table](#)

Other diagnostics: [detect_intrusions\(\)](#), [ese\(\)](#), [pdi\(\)](#), [phase_diagnostic_table\(\)](#), [predict_phase\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
head(as_phase_table(fit))
```

as_plotly	<i>Convert a ggplot to an interactive plotly widget</i>
-----------	---

Description

Wraps [ggplotly](#) with archaeological tooltips showing find ID, context, phase probability, dating, class, entropy, and energy.

Usage

```
as_plotly(gg, tooltip = "text", ...)
```

Arguments

gg	A ggplot object produced by any gg_* function.
tooltip	Character vector of aesthetics to show. Defaults to "text" which displays the enriched archaeological tooltip.
...	Additional arguments passed to ggplotly .

Value

A plotly htmlwidget object.

See Also

[gg_phasefield](#), [gg_entropy](#), [gg_energy](#), [gg_intrusions](#)

Other plotting: [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1)
fit <- fit_sef(x, k = 3)
if (requireNamespace("ggplot2", quietly = TRUE) &&
    requireNamespace("plotly", quietly = TRUE)) {
  p <- as_plotly(gg_phasefield(fit))
}
```

as_sf_links

Export high-SEI links as an sf LINESTRING layer

Description

Extracts the strongest pairwise SEI connections as sf line geometries.

Usage

```
as_sf_links(object, quantile_threshold = 0.9, crs = NA_integer_)
```

Arguments

object	A sef_fit object.
quantile_threshold	Quantile for retaining strongest links (default: 0.9).
crs	CRS for the output geometry.

Value

An sf object with columns from, to, sei, and geometry.

See Also

[as_sf_phase](#), [sei_matrix](#)

Other GIS: [as_sf_phase\(\)](#)

Examples

```
if (requireNamespace("sf", quietly = TRUE)) {
  x <- archaeo_sim(n = 60, k = 2, seed = 1)
  fit <- fit_sef(x, k = 2)
  links <- as_sf_links(fit)
}
```

as_sf_phase

Convert a fitted model to an sf point layer

Description

Creates an sf point object with phase assignments and diagnostics for use in QGIS or spatial analysis.

Usage

```
as_sf_phase(object, crs = NA_integer_, dims = c("XY", "XYZ"))
```

Arguments

object	A sef_fit object.
crs	CRS passed to st_as_sf .
dims	Either "XY" or "XYZ".

Value

An sf object.

See Also

[as_sf_links](#), [phase_diagnostic_table](#)

Other GIS: [as_sf_links\(\)](#)

Examples

```

if (requireNamespace("sf", quietly = TRUE)) {
  x <- archaeo_sim(n = 60, k = 2, seed = 1)
  fit <- fit_sef(x, k = 2)
  pts <- as_sf_phase(fit)
}

```

bootstrap_sef

*Bootstrap confidence intervals for SEF diagnostics***Description**

Resamples the data with replacement, refits the model, and computes the distribution of key statistics (PDI, mean entropy, mean energy, ARI if true labels provided). Returns percentile confidence intervals.

Usage

```

bootstrap_sef(
  object,
  n_boot = 100,
  conf = 0.95,
  true_labels = NULL,
  verbose = TRUE
)

```

Arguments

object	A <code>sef_fit</code> object (used to extract fitting parameters).
n_boot	Number of bootstrap replicates (default: 100).
conf	Confidence level (default: 0.95).
true_labels	Optional integer vector of true phase labels for ARI.
verbose	Print progress (default: TRUE).

Value

A data.frame with columns `statistic`, `estimate`, `lower`, `upper`, `se`.

See Also

[fit_sef](#), [pdi](#), [adjusted_rand_index](#)

Other validation: [adjusted_rand_index\(\)](#), [confusion_matrix\(\)](#), [cv_sef\(\)](#), [optimize_weights\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2, seed = 1)
bootstrap_sef(fit, n_boot = 20)
```

`compare_k`*Compare multiple candidate phase counts*

Description

Fits the SEF model for each value of K and returns a summary table with BIC, PDI, entropy, energy, and other diagnostics.

Usage

```
compare_k(data, k_values = 2:6, ...)
```

Arguments

<code>data</code>	Input data.frame.
<code>k_values</code>	Integer vector of candidate phase counts.
<code>...</code>	Additional arguments passed to fit_sef .

Value

A data.frame with one row per K value.

See Also

[fit_sef](#), [gg_compare_k](#)

Other fitting: [fit_sef\(\)](#), [reorder_phases\(\)](#), [sef_summary\(\)](#)

Examples

```
x <- archaeo_sim(n = 100, k = 3, seed = 1)
ck <- compare_k(x, k_values = 2:4)
print(ck)
```

confusion_matrix	<i>Confusion matrix between estimated and true phases</i>
------------------	---

Description

Cross-tabulates estimated phase assignments against known true labels. Phases are reordered to maximise diagonal agreement (Hungarian matching).

Usage

```
confusion_matrix(object, true_labels)
```

Arguments

object	A <code>sef_fit</code> object, or an integer vector of predicted labels.
true_labels	Integer vector of known phase assignments.

Value

A matrix with estimated phases as rows and true phases as columns.

See Also

[adjusted_rand_index](#)

Other validation: [adjusted_rand_index\(\)](#), [bootstrap_sef\(\)](#), [cv_sef\(\)](#), [optimize_weights\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1, mixing = 0.05)
fit <- fit_sef(x, k = 3, seed = 1)
confusion_matrix(fit, x$true_phase)
```

cv_sef	<i>K-fold cross-validation for SEF model</i>
--------	--

Description

Splits the data into folds, fits on training folds, and evaluates log-likelihood on the held-out fold. Useful for comparing different K values or weight configurations.

Usage

```
cv_sef(data, k_values = 2:6, n_folds = 5, seed = 1, ...)
```

Arguments

data	A data.frame with archaeological find data.
k_values	Integer vector of candidate phase counts.
n_folds	Number of cross-validation folds (default: 5).
seed	Random seed for fold assignment.
...	Additional arguments passed to fit_sef .

Value

A data.frame with columns k, fold, train_loglik, test_loglik, train_pdi.

See Also

[compare_k](#), [fit_sef](#)

Other validation: [adjusted_rand_index\(\)](#), [bootstrap_sef\(\)](#), [confusion_matrix\(\)](#), [optimize_weights\(\)](#)

Examples

```
x <- archaeo_sim(n = 100, k = 3, seed = 1)
cv <- cv_sef(x, k_values = 2:4, n_folds = 3)
# Mean test log-likelihood per K
aggregate(test_loglik ~ k, data = cv, FUN = mean)
```

demo_compressed

Demo dataset: compressed palimpsest

Description

Simulated dataset with 250 artefacts, 4 phases, and 50% mixing. Represents a heavily disturbed deposit where phases are difficult to separate.

Usage

```
demo_compressed
```

Format

A data.frame with 250 rows and 10 columns. See [demo_easy](#) for column descriptions.

Examples

```
data(demo_compressed)

ck <- compare_k(demo_compressed, k_values = 2:6)
print(ck)
```

`demo_easy`*Demo dataset: well-separated phases*

Description

Simulated dataset with 150 artefacts, 3 phases, and 5% mixing. Represents a site where occupation phases are clearly distinguishable.

Usage`demo_easy`**Format**

A data.frame with 150 rows and 10 columns:

id Artefact identifier

x, y Planimetric coordinates (metres)

z Depth (metres)

context Stratigraphic unit label

date_min, date_max Chronological interval (CE)

class Cultural class (ceramic, lithic, bone, metal)

taf_score Taphonomic disturbance score (0-1)

true_phase Known phase assignment (for validation)

Examples

```
data(demo_easy)
fit <- fit_sef(demo_easy, k = 3)
plot_phasefield(fit)
```

`demo_moderate`*Demo dataset: moderate palimpsest*

Description

Simulated dataset with 200 artefacts, 3 phases, and 30% mixing. Represents a site with significant but resolvable depositional mixing.

Usage`demo_moderate`

Format

A data.frame with 200 rows and 10 columns. See [demo_easy](#) for column descriptions.

Examples

```
data(demo_moderate)
fit <- fit_sef(demo_moderate, k = 3, tafonomy = "taf_score", context = "context")
summary(fit)
```

detect_intrusions	<i>Detect potentially intrusive observations</i>
-------------------	--

Description

Combines rescaled entropy, energy, and inverse local SEI into a composite intrusion probability score.

Usage

```
detect_intrusions(object)
```

Arguments

object A sef_fit object.

Value

A data.frame with columns id and intrusion_prob.

See Also

[gg_intrusions](#), [fit_sef](#)

Other diagnostics: [as_phase_table\(\)](#), [ese\(\)](#), [pdi\(\)](#), [phase_diagnostic_table\(\)](#), [predict_phase\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
di <- detect_intrusions(fit)
head(di[order(di$intrusion_prob, decreasing = TRUE), ])
```

ese

Compute Excavation Stratigraphic Energy

Description

Measures local depositional disruption for each find by summing weighted dissimilarities with neighbours.

Usage

```
ese(  
  data,  
  coords = c("x", "y", "z"),  
  chrono = c("date_min", "date_max"),  
  class_col = "class",  
  beta = c(1, 1, 1, 1),  
  neighbourhood = NULL  
)
```

Arguments

data	Input data.frame.
coords	Character vector of coordinate column names.
chrono	Character vector with minimum and maximum dating columns.
class_col	Class column name.
beta	Numeric vector of length 4: weights for spatial, vertical, temporal, and class mismatch.
neighbourhood	Maximum XY distance for neighbour inclusion. When NULL, all observations contribute.

Value

A numeric vector of local energy values.

See Also

[fit_sef](#), [gg_energy](#)

Other diagnostics: [as_phase_table\(\)](#), [detect_intrusions\(\)](#), [pdi\(\)](#), [phase_diagnostic_table\(\)](#), [predict_phase\(\)](#)

Examples

```
x <- archaeo_sim(n = 30, k = 2, seed = 1)  
e <- ese(x)  
summary(e)
```

export_results	<i>Export all results to files</i>
----------------	------------------------------------

Description

Writes phase assignments, intrusion scores, US summary, and model summary to CSV files in the specified directory.

Usage

```
export_results(object, dir, format = "csv", prefix = "palimpsestr")
```

Arguments

object	A sef_fit object.
dir	Output directory (created if it does not exist). Must be specified explicitly; consider using <code>tempdir()</code> for temporary output.
format	Export format: "csv" (default).
prefix	File name prefix (default: "palimpsestr").

Value

Invisibly returns a character vector of written file paths.

See Also

[us_summary_table](#), [as_phase_table](#)

Other export: [phase_transition_matrix\(\)](#), [us_summary_table\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2, context = "context")
export_results(fit, dir = tempdir())
```

 fit_sef

Fit the Stratigraphic Entanglement Field model

Description

Estimates latent depositional phases from spatial, stratigraphic, chronological, and cultural evidence using diagonal Gaussian mixture EM.

Usage

```
fit_sef(
  data,
  coords = c("x", "y", "z"),
  chrono = c("date_min", "date_max"),
  class = "class",
  tafonomy = NULL,
  context = NULL,
  harris = NULL,
  k = 3,
  weights = c(ws = 1, wz = 1, wt = 1, wc = 1),
  seed = 1,
  em_iter = 100,
  em_tol = 1e-05,
  n_init = 1,
  chrono_precision = FALSE,
  taf_as_feature = FALSE,
  residuality = FALSE,
  class_scale = FALSE,
  subclass = NULL
)
```

Arguments

data	A data.frame with archaeological find data.
coords	Character vector of length 3 naming the x, y, z coordinate columns.
chrono	Character vector of length 2 naming the min and max dating columns.
class	Character scalar naming the material class column.
tafonomy	Optional column name for taphonomic disturbance scores (0–1).
context	Optional column name for stratigraphic unit labels.
harris	Optional $n \times n$ matrix of pairwise stratigraphic penalties.
k	Integer number of phases to estimate.
weights	Named numeric vector with components ws, wz, wt, wc.
seed	Random seed for reproducibility.
em_iter	Maximum number of EM iterations (default: 100).

em_tol	Convergence tolerance on the log-likelihood.
n_init	Number of random initialisations. The run with the highest log-likelihood is retained (default: 1).
chrono_precision	Logical. If TRUE, add 1/tspan as a feature giving higher weight to precisely dated finds (default: FALSE).
taf_as_feature	Logical. If TRUE and tafonomy is provided, include taf_score as an additional feature dimension (default: FALSE).
residuality	Logical. If TRUE and context is provided, add a residuality score measuring chronological mismatch between each find and its context mean (default: FALSE).
class_scale	Logical. If TRUE, scale one-hot class columns by $1/\sqrt{n_classes}$ so their total energy is comparable to a single numeric feature (default: FALSE).
subclass	Optional column name for sub-class labels. If provided, used instead of class for one-hot encoding in the feature matrix.

Value

An S3 object of class `sef_fit`.

See Also

[archaeo_sim](#), [compare_k](#), [pdi](#), [detect_intrusions](#)

Other fitting: [compare_k\(\)](#), [reorder_phases\(\)](#), [sef_summary\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
print(fit)
```

```
x <- archaeo_sim(n = 150, k = 3, seed = 42)
fit <- fit_sef(x, k = 3, tafonomy = "taf_score", context = "context")
summary(fit)
```

gg_bootstrap

Bootstrap confidence interval plot

Description

Visualises bootstrap confidence intervals for key SEF diagnostics (PDI, entropy, energy, log-likelihood, and optionally ARI).

Usage

```
gg_bootstrap(bs_result)
```

Arguments

bs_result Data.frame returned by [bootstrap_sef](#).

Value

A ggplot object.

See Also

[bootstrap_sef](#)

Other plotting: [as_plotly\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_compare_k

Phase count selection diagnostics

Description

Three-panel plot showing BIC, PDI, and Mean Entropy for different K values.

Usage

```
gg_compare_k(ck)
```

Arguments

ck Data.frame from [compare_k\(\)](#).

Value

A ggplot object.

See Also

[compare_k](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_confusion	<i>Confusion matrix heatmap</i>
--------------	---------------------------------

Description

Plots a heatmap of the confusion matrix between estimated and known true phase assignments.

Usage

```
gg_confusion(object, true_labels)
```

Arguments

object	A <code>sef_fit</code> object.
true_labels	Integer vector of known true phase assignments.

Value

A ggplot object.

See Also

[confusion_matrix](#), [adjusted_rand_index](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
if (requireNamespace("ggplot2", quietly = TRUE)) {  
  x <- archaeo_sim(n = 80, k = 3, seed = 1, mixing = 0.05)  
  fit <- fit_sef(x, k = 3, seed = 1)  
  gg_confusion(fit, x$true_phase)  
}
```

gg_convergence	<i>EM convergence trace</i>
----------------	-----------------------------

Description

Plots the log-likelihood at each EM iteration to verify convergence.

Usage

```
gg_convergence(object)
```

Arguments

object A `sef_fit` object.

Value

A ggplot object.

See Also

[fit_sef](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
if (requireNamespace("ggplot2", quietly = TRUE)) {  
  x <- archaeo_sim(n = 80, k = 3, seed = 1)  
  fit <- fit_sef(x, k = 3)  
  gg_convergence(fit)  
}
```

gg_cv

Cross-validation diagnostic plot

Description

Plots mean test and training log-likelihood across K values from `cv_sef` output.

Usage

```
gg_cv(cv_result)
```

Arguments

cv_result Data.frame returned by `cv_sef`.

Value

A ggplot object.

See Also

[cv_sef](#), [gg_compare_k](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_energy	<i>Excavation Stratigraphic Energy map</i>
-----------	--

Description

Shows ESE values across space. High energy indicates zones where neighbouring artefacts are dissimilar (depositional disruption).

Usage

```
gg_energy(object, xlabel = "Easting (m)", ylabel = "Northing (m)")
```

Arguments

object	A sef_fit object.
xlabel, ylabel	Axis labels.

Value

A ggplot object.

See Also

[gg_entropy](#), [as_plotly](#) for interactive version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_entropy	<i>Spatial entropy map</i>
------------	----------------------------

Description

Shows Shannon entropy of phase probabilities across the excavation area. High entropy = uncertain phase assignment (possible palimpsest zone).

Usage

```
gg_entropy(object, xlabel = "Easting (m)", ylabel = "Northing (m)")
```

Arguments

object	A sef_fit object.
xlabel, ylabel	Axis labels.

Value

A ggplot object.

See Also

[plot_entropy](#) for base R version, [as_plotly](#) for interactive version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_intrusions

Intrusion probability map

Description

Maps the probability that each find is an intrusion (displaced or redeposited). Top suspects are circled and labelled.

Usage

```
gg_intrusions(
  object,
  top_n = 5,
  xlabel = "Easting (m)",
  ylabel = "Northing (m)"
)
```

Arguments

`object` A `sef_fit` object.
`top_n` Number of top intrusions to label.
`xlabel, ylabel` Axis labels.

Value

A ggplot object.

See Also

[detect_intrusions](#), [as_plotly](#) for interactive version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

`gg_map`*Overlay SEF results on excavation plan geometries*

Description

Displays SEF analysis results directly on the excavation plan. Polygons represent stratigraphic units; points represent individual finds.

Usage

```
gg_map(  
  object,  
  geometries,  
  layer = c("phase", "entropy", "energy", "intrusion"),  
  show_labels = TRUE,  
  show_points = TRUE,  
  xlabel = NULL,  
  ylabel = NULL  
)
```

Arguments

<code>object</code>	A <code>sef_fit</code> object.
<code>geometries</code>	An <code>sf</code> object with excavation plan polygons. Must contain a <code>us</code> column matching context values in the data.
<code>layer</code>	What to display: "phase", "entropy", "energy", or "intrusion".
<code>show_labels</code>	Show US labels on polygons.
<code>show_points</code>	Overlay individual finds as points.
<code>xlabel, ylabel</code>	Axis labels (default: NULL = use CRS units).

Value

A `ggplot` object.

See Also

[load_geometries](#), [as_sf_phase](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_phase_profile *Vertical phase profile*

Description

Plots finds along the depth (z) axis, coloured by phase assignment, to visualise the stratigraphic ordering of phases.

Usage

```
gg_phase_profile(object)
```

Arguments

object A sef_fit object.

Value

A ggplot object.

See Also

[phase_transition_matrix](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
if (requireNamespace("ggplot2", quietly = TRUE)) {
  x <- archaeo_sim(n = 80, k = 3, seed = 1)
  fit <- fit_sef(x, k = 3)
  gg_phase_profile(fit)
}
```

gg_phasefield *Phase assignment map*

Description

Plots artefact positions coloured by dominant phase, with point size proportional to assignment confidence.

Usage

```
gg_phasefield(object, xlabel = "Easting (m)", ylabel = "Northing (m)")
```

Arguments

object A `sef_fit` object.
xlabel, ylabel Axis labels (default: "Easting (m)" / "Northing (m)").

Value

A ggplot object.

See Also

[plot_phasefield](#) for base R version, [as_plotly](#) for interactive version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

gg_weights	<i>Weight sensitivity heatmap</i>
------------	-----------------------------------

Description

Plots the mean test log-likelihood across weight configurations from [optimize_weights](#) output.

Usage

```
gg_weights(opt_result)
```

Arguments

opt_result List returned by [optimize_weights](#).

Value

A ggplot object.

See Also

[optimize_weights](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

`harris_from_contexts` *Generate stratigraphic penalty matrix from context depth ordering*

Description

Infers vertical ordering between stratigraphic units from the mean depth of finds in each context, and builds a penalty matrix that discourages finds from different vertical zones being assigned to the same phase.

Usage

```
harris_from_contexts(  
  data,  
  z_col = "z",  
  context_col = "context",  
  penalty_scale = 0.5  
)
```

Arguments

<code>data</code>	A data.frame with find data.
<code>z_col</code>	Name of the depth column.
<code>context_col</code>	Name of the context column.
<code>penalty_scale</code>	Penalty magnitude for cross-context assignments.

Value

An $n \times n$ symmetric penalty matrix.

See Also

[fit_sef](#), [read_harris](#)

Other harris: [read_harris\(\)](#), [validate_phases_harris\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)  
H <- harris_from_contexts(x, z_col = "z", context_col = "context")  
dim(H)
```

launch_app	<i>Launch the palimpsestr Shiny Dashboard</i>
------------	---

Description

Opens an interactive Shiny application for loading data, fitting the SEF model, and exploring results through plots, tables, and maps.

Usage

```
launch_app()
```

Details

The app requires the following packages (installed but not imported by palimpsestr): shiny, shinydashboard, DT. For interactive plots install plotly. For database connections install DBI plus RSQLite or RPostgres. For GIS maps install sf.

Value

Launches the Shiny app (does not return).

Examples

```
## Not run:  
launch_app()  
  
## End(Not run)
```

load_geometries	<i>Load excavation geometries from file or database</i>
-----------------	---

Description

Reads stratigraphic unit polygons from Shapefile, GeoPackage, GeoJSON, or a PostGIS database for use with [gg_map](#).

Usage

```
load_geometries(  
  source,  
  layer = NULL,  
  query = NULL,  
  us_column = "us",  
  crs = NULL  
)
```

Arguments

source	File path or DBIConnection.
layer	Layer name for multi-layer sources.
query	SQL query for database connections.
us_column	Column containing US/context identifiers.
crs	Target CRS for reprojection (optional).

Value

An sf object with a us column.

See Also

[gg_map](#), [read_db](#)

Other data-import: [read_db\(\)](#)

local_sei	<i>Compute local SEI values</i>
-----------	---------------------------------

Description

Aggregates the SEI matrix by row, yielding a per-observation measure of total depositional coherence with all other finds.

Usage

```
local_sei(sei_mat)
```

Arguments

sei_mat	A symmetric SEI matrix from sei_matrix .
---------	--

Value

A numeric vector of length `nrow(sei_mat)`.

See Also

[sei_matrix](#)

Other SEI: [sei_matrix\(\)](#), [sei_sparse\(\)](#)

Examples

```
x <- archaeo_sim(n = 30, k = 2, seed = 1)
S <- sei_matrix(x)
lsei <- local_sei(S)
summary(lsei)
```

optimize_weights *Estimate optimal SEI weights via cross-validation*

Description

Tests a grid of weight configurations and selects the one that maximises the mean held-out log-likelihood across folds. This provides a data-driven alternative to manual weight specification.

Usage

```
optimize_weights(  
  data,  
  k,  
  weight_grid = NULL,  
  n_folds = 3,  
  seed = 1,  
  verbose = TRUE,  
  ...  
)
```

Arguments

data	A data.frame with archaeological find data.
k	Number of phases.
weight_grid	A data.frame with columns ws, wz, wt, wc. If NULL, a default grid is used.
n_folds	Number of cross-validation folds (default: 3).
seed	Random seed.
verbose	Print progress (default: TRUE).
...	Additional arguments passed to fit_sef .

Value

A list with components:

best_weights Named numeric vector of optimal weights.

results Data.frame with all tested configurations and their scores.

See Also

[fit_sef](#), [cv_sef](#)

Other validation: [adjusted_rand_index\(\)](#), [bootstrap_sef\(\)](#), [confusion_matrix\(\)](#), [cv_sef\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1)
opt <- optimize_weights(x, k = 3, n_folds = 3)
opt$best_weights
```

pdi

Compute Palimpsest Dissolution Index

Description

Measures global phase separability as $1 - \bar{H}/\log(K)$. Values close to 1 indicate well-separated phases; values near 0 indicate a compressed palimpsest.

Usage

```
pdi(object)
```

Arguments

object A `sef_fit` object.

Value

A single numeric value between 0 and 1.

See Also

[fit_sef](#), [compare_k](#)

Other diagnostics: [as_phase_table\(\)](#), [detect_intrusions\(\)](#), [ese\(\)](#), [phase_diagnostic_table\(\)](#), [predict_phase\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
pdi(fit)
```

`phase_diagnostic_table`*Return a compact diagnostic table*

Description

Combines the input data with dominant phase, phase probabilities, entropy, local SEI, and energy in a single data.frame.

Usage

```
phase_diagnostic_table(object)
```

Arguments

`object` A `sef_fit` object.

Value

A data.frame with all input columns plus diagnostics.

See Also

[as_phase_table](#), [as_sf_phase](#)

Other diagnostics: [as_phase_table\(\)](#), [detect_intrusions\(\)](#), [ese\(\)](#), [pdi\(\)](#), [predict_phase\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
pdt <- phase_diagnostic_table(fit)
names(pdt)
```

`phase_transition_matrix`*Phase vertical transition matrix*

Description

Computes how often finds from phase *i* are found directly above finds from phase *j* in the vertical sequence, revealing the stratigraphic ordering of phases.

Usage

```
phase_transition_matrix(object)
```

Arguments

object A sef_fit object.

Value

A $K \times K$ matrix where entry $[i, j]$ counts transitions from phase i (above) to phase j (below).

See Also

[us_summary_table](#)

Other export: [export_results\(\)](#), [us_summary_table\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1)
fit <- fit_sef(x, k = 3)
phase_transition_matrix(fit)
```

plot_energy

Plot local Excavation Stratigraphic Energy (base R)

Description

Plot local Excavation Stratigraphic Energy (base R)

Usage

```
plot_energy(object)
```

Arguments

object A sef_fit object.

Value

Invisibly returns the object.

See Also

[gg_energy](#) for the ggplot2 version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
plot_energy(fit)
```

plot_entropy	<i>Plot entropy across space (base R)</i>
--------------	---

Description

Plot entropy across space (base R)

Usage

```
plot_entropy(object)
```

Arguments

object A `sef_fit` object.

Value

Invisibly returns the object.

See Also

[gg_entropy](#) for the ggplot2/plotly version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_phasefield\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
plot_entropy(fit)
```

plot_phasefield	<i>Plot dominant phase assignment (base R)</i>
-----------------	--

Description

Plot dominant phase assignment (base R)

Usage

```
plot_phasefield(object)
```

Arguments

object A `sef_fit` object.

Value

Invisibly returns the object.

See Also

[gg_phasefield](#) for the ggplot2/plotly version.

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_sei_profile\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
plot_phasefield(fit)
```

plot_sei_profile	<i>Plot ordered SEI profile (base R)</i>
------------------	--

Description

Plot ordered SEI profile (base R)

Usage

```
plot_sei_profile(object)
```

Arguments

object A sef_fit object.

Value

Invisibly returns the object.

See Also

[sei_matrix](#), [local_sei](#)

Other plotting: [as_plotly\(\)](#), [gg_bootstrap\(\)](#), [gg_compare_k\(\)](#), [gg_confusion\(\)](#), [gg_convergence\(\)](#), [gg_cv\(\)](#), [gg_energy\(\)](#), [gg_entropy\(\)](#), [gg_intrusions\(\)](#), [gg_map\(\)](#), [gg_phase_profile\(\)](#), [gg_phasefield\(\)](#), [gg_weights\(\)](#), [plot_energy\(\)](#), [plot_entropy\(\)](#), [plot_phasefield\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
plot_sei_profile(fit)
```

predict_phase	<i>Predict phase probabilities</i>
---------------	------------------------------------

Description

Convenience alias for [as_phase_table](#).

Usage

```
predict_phase(object)
```

Arguments

object A `sef_fit` object.

Value

A `data.frame` with probabilities and diagnostics.

See Also

[as_phase_table](#)

Other diagnostics: [as_phase_table\(\)](#), [detect_intrusions\(\)](#), [ese\(\)](#), [pdi\(\)](#), [phase_diagnostic_table\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
head(predict_phase(fit))
```

read_db	<i>Read archaeological data from a SQL database</i>
---------	---

Description

Loads find data from any DBI-compatible database and maps columns to the format expected by [fit_sef](#).

Usage

```
read_db(
  conn,
  query = NULL,
  table = NULL,
  col_id = "id",
  col_x = "x",
  col_y = "y",
  col_z = "z",
  col_context = "context",
  col_date_min = "date_min",
  col_date_max = "date_max",
  col_class = "class",
  col_taf = NULL,
  schema = "custom",
  sito = NULL
)
```

Arguments

conn	A DBIConnection object.
query	SQL query returning data with the required columns.
table	Table name (alternative to query).
col_id, col_x, col_y, col_z, col_context, col_date_min, col_date_max, col_class, col_taf	Column name mappings.
schema	Use "pyarchinit" for automatic PyArchInit mapping.
sito	Site filter for PyArchInit schema.

Value

A data.frame ready for [fit_sef](#).

See Also

[fit_sef](#), [load_geometries](#)

Other data-import: [load_geometries\(\)](#)

read_harris

Read Harris Matrix from CSV edge list

Description

Reads a CSV file with columns from, to, and optionally weight, and converts it to an $n \times n$ penalty matrix aligned with the find-level data.

Usage

```
read_harris(file, contexts, default_weight = 1)
```

Arguments

`file` Path to CSV with columns from, to, and optionally weight.
`contexts` Character vector of context labels for each find (length = number of finds).
`default_weight` Weight for edges without an explicit weight.

Value

An $n \times n$ penalty matrix.

See Also

[harris_from_contexts](#), [fit_sef](#)

Other harris: [harris_from_contexts\(\)](#), [validate_phases_harris\(\)](#)

reorder_phases	<i>Reorder phases by mean depth</i>
----------------	-------------------------------------

Description

Relabels phases so that phase 1 corresponds to the deepest (oldest) stratum and phase K to the shallowest (most recent). This ensures consistent, interpretable phase numbering across different runs.

Usage

```
reorder_phases(object)
```

Arguments

`object` A `sef_fit` object.

Value

A `sef_fit` object with reordered phases.

See Also

[fit_sef](#)

Other fitting: [compare_k\(\)](#), [fit_sef\(\)](#), [sef_summary\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1)
fit <- fit_sef(x, k = 3)
fit <- reorder_phases(fit)
```

report_sef	<i>Generate a textual interpretive report for a SEF model</i>
------------	---

Description

Produces a structured Markdown report covering phase composition, intrusion detection, stratigraphic unit purity, and recommendations. Available in English and Italian.

Usage

```
report_sef(object, lang = c("en", "it"), file = NULL)
```

Arguments

object	A sef_fit object.
lang	Language: "en" (English) or "it" (Italian).
file	Optional file path to save the report.

Value

Character string with the report text (invisibly).

See Also

[fit_sef](#), [sef_summary](#)

Examples

```
x <- archaeo_sim(n = 100, k = 3, seed = 1)
fit <- fit_sef(x, k = 3)
report_sef(fit, lang = "en")
```

sef_summary	<i>Compact summary for a fitted SEF model</i>
-------------	---

Description

Returns a named list with global diagnostics and phase counts.

Usage

```
sef_summary(object)
```

Arguments

object	A sef_fit object.
--------	-------------------

Value

A named list.

See Also

[fit_sef](#), [pdi](#)

Other fitting: [compare_k\(\)](#), [fit_sef\(\)](#), [reorder_phases\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 2, seed = 1)
fit <- fit_sef(x, k = 2)
sef_summary(fit)
```

 sei_matrix

Compute the Stratigraphic Entanglement Index matrix

Description

Builds an $n \times n$ symmetric matrix quantifying pairwise depositional coherence from spatial, vertical, temporal, and cultural evidence.

Usage

```
sei_matrix(
  data,
  coords = c("x", "y", "z"),
  chrono = c("date_min", "date_max"),
  class_col = "class",
  weights = c(ws = 1, wz = 1, wt = 1, wc = 1),
  eps = 1e-09,
  z_floor = 0.25,
  max_dist = NULL
)
```

Arguments

data	Input data.frame.
coords	Character vector of coordinate column names (x, y, z).
chrono	Character vector with minimum and maximum dating columns.
class_col	Class column name.
weights	Named numeric vector with components ws, wz, wt, wc. Each component is normalised to [0, 1] before weighting, so the weights represent relative importance.
eps	Small value to avoid division by zero in spatial distance.
z_floor	Minimum vertical denominator.

`max_dist` Maximum spatial distance for pair inclusion. When NULL (default), all pairs are computed. For large datasets ($n > 2000$), setting this to a reasonable neighbourhood radius dramatically reduces memory and computation time. The result is a sparse-like matrix with zeros for distant pairs.

Value

A symmetric numeric matrix with zero diagonal.

Note

SEI values are normalised within each dataset. Absolute SEI values are **not directly comparable** across different excavations or datasets of different sizes. Use SEI for within-dataset ranking and relative comparisons only.

See Also

[local_sei](#), [fit_sef](#)

Other SEI: [local_sei\(\)](#), [sei_sparse\(\)](#)

Examples

```
x <- archaeo_sim(n = 30, k = 2, seed = 1)
S <- sei_matrix(x)
dim(S)
```

`sei_sparse`

Compute SEI matrix with automatic sparsification

Description

Wrapper around [sei_matrix](#) that automatically sets `max_dist` when the dataset exceeds a size threshold, using the 25th percentile of pairwise distances as the cutoff.

Usage

```
sei_sparse(
  data,
  coords = c("x", "y", "z"),
  chrono = c("date_min", "date_max"),
  class_col = "class",
  weights = c(ws = 1, wz = 1, wt = 1, wc = 1),
  eps = 1e-09,
  z_floor = 0.25,
  n_threshold = 1000
)
```

Arguments

data	Input data.frame.
coords	Character vector of coordinate column names (x, y, z).
chrono	Character vector with minimum and maximum dating columns.
class_col	Class column name.
weights	Named numeric vector with components ws, wz, wt, wc. Each component is normalised to [0, 1] before weighting, so the weights represent relative importance.
eps	Small value to avoid division by zero in spatial distance.
z_floor	Minimum vertical denominator.
n_threshold	Datasets larger than this trigger sparse mode (default: 1000).

Value

A numeric matrix (same as sei_matrix).

See Also

[sei_matrix](#)

Other SEI: [local_sei\(\)](#), [sei_matrix\(\)](#)

Examples

```
x <- archaeo_sim(n = 50, k = 2, seed = 1)
S <- sei_sparse(x)
```

summary.sef_fit	<i>Summarise a fitted SEF model</i>
-----------------	-------------------------------------

Description

Summarise a fitted SEF model

Usage

```
## S3 method for class 'sef_fit'
summary(object, ...)
```

Arguments

object	A sef_fit object.
...	Ignored.

Value

A named list.

us_summary_table	<i>Summary table per stratigraphic unit</i>
------------------	---

Description

Aggregates finds by context (US), reporting the dominant phase, purity (proportion of finds in dominant phase), mean entropy, mean energy, and intrusion count.

Usage

```
us_summary_table(object)
```

Arguments

object A sef_fit object.

Value

A data.frame with one row per stratigraphic unit.

See Also

[export_results](#), [phase_diagnostic_table](#)

Other export: [export_results\(\)](#), [phase_transition_matrix\(\)](#)

Examples

```
x <- archaeo_sim(n = 80, k = 3, seed = 1)
fit <- fit_sef(x, k = 3, context = "context")
us_summary_table(fit)
```

validate_phases_harris

Validate phase assignments against stratigraphic ordering

Description

Checks whether the estimated phases follow the expected vertical ordering within each stratigraphic unit pair.

Usage

```
validate_phases_harris(object)
```

Arguments

object A sef_fit object.

Value

A data.frame with one row per context pair, indicating whether the dominant phase ordering is consistent with depth.

See Also

[harris_from_contexts](#), [us_summary_table](#)

Other harris: [harris_from_contexts\(\)](#), [read_harris\(\)](#)

Examples

```
x <- archaeo_sim(n = 60, k = 3, seed = 1)
fit <- fit_sef(x, k = 3, context = "context")
validate_phases_harris(fit)
```

villa_romana

Poggio Gramignano Roman Villa excavation dataset

Description

Real archaeological dataset from the site of Poggio Gramignano (Lugnano in Teverina, TR, Italy), a multi-period Roman villa with an annexed Late Antique infant cemetery. The dataset contains 615 inventoried finds from 54 stratigraphic units spanning from the Pre-Roman period (7th–4th c. BCE) through the Late Antique (5th–6th c. CE).

Usage

villa_romana

Format

A data.frame with 615 rows and 9 variables:

id Unique find identifier (VRPG_0001 to VRPG_0615)

x Easting coordinate (metres, EPSG:3004)

y Northing coordinate (metres, EPSG:3004)

z Mean elevation of the stratigraphic unit (metres a.s.l.)

context Stratigraphic unit label (e.g. US_76, US_117)

date_min Start of chronological interval (BCE as negative, CE as positive)

date_max End of chronological interval

class Material class (e.g. Reperto Ceramico, Reperto Anforaceo)

taf_score Taphonomic integrity score (0 = fully disturbed, 1 = pristine in situ)

Details

Chronological data derives from the site periodisation. Coordinates are US centroids in Monte Mario / Italy zone 2 (EPSG:3004). Elevation (z) is the mean of recorded quotes per US. Taphonomic scores were assigned by the excavator based on depositional context (primary, secondary, redeposition, backfill, disturbance).

Source

Data from the pyArchInit database of Poggio Gramignano (VRPG 3004). Taphonomic scores compiled by Roberto Montagnetti.

Examples

```
data(villa_romana)
str(villa_romana)
table(villa_romana$class)
```

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