

Package ‘smile’

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Title Spatial Misalignment: Interpolation, Linkage, and Estimation

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Description Provides functions to estimate, predict and interpolate areal data. For estimation and prediction we assume areal data is an average of an underlying continuous spatial process as in Moraga et al. (2017) <[doi:10.1016/j.spasta.2017.04.006](https://doi.org/10.1016/j.spasta.2017.04.006)>, Johnson et al. (2020) <[doi:10.1186/s12942-020-00200-w](https://doi.org/10.1186/s12942-020-00200-w)>, and Wilson and Wakefield (2020) <[doi:10.1093/biostatistics/kxy041](https://doi.org/10.1093/biostatistics/kxy041)>. The interpolation methodology is (mostly) based on Goodchild and Lam (1980, ISSN:01652273).

License GPL-3

Encoding UTF-8

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SystemRequirements GDAL (>= 2.0.1), GEOS (>= 3.4.0), PROJ (>= 4.8.0)

LinkingTo Rcpp, RcppEigen

Imports numDeriv, Rcpp, sf, mvtnorm, stats, parallel, Matrix

Depends R (>= 4.1.0)

URL <https://lcgodoy.me/smile/>, <https://github.com/lcgodoy/smile/>

BugReports <https://github.com/lcgodoy/smile/issues/>

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AI	<i>Areal Interpolation</i>
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Description

This function estimates variables observed at a "source" region into a "target" region. "Source" and "target" regions represent two different ways to divide a city, for example. For more details, see <https://lsgodoy.me/smile/articles/sai.html>.

Usage

```
ai(source, target, vars)
```

```
ai_var(source, target, vars, vars_var, sc_vars = FALSE, var_method = "CS")
```

Arguments

source	a sf object - source spatial data.
target	a sf object - target spatial data.
vars	a character representing the variables (observed at the source) to be estimated at the target data.
vars_var	a scalar of type character representing the name of the variable in the source dataset that stores the variances of the variable to be estimated at the target data.
sc_vars	boolean indicating whether vars should be scaled by its observed variance (if available).
var_method	a character representing the method to approximate the variance of the AI estimates. Possible values are "CS" (Cauchy-Schwartz) or "MI" (Moran's I).

Value

the target (of type sf) with estimates of the variables observed at the source data.

find_phi	<i>Find phi parameter for the Exponential spatial auto-correlation function</i>
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Description

Function designed to find the phi parameter such that the correlation between points within a given distance d is at most a given value.

Usage

```
find_phi(
  d,
  nu,
  kappa,
  mu2,
  family = "matern",
  range = c(1e-04, 1000),
  cut = 0.05
)
```

Arguments

d	maximum distance for spatial dependence equal to cut.
nu	smoothness parameter associated with the Matern cov. function.
kappa	one of the smoothness parameters associated with the Generalized Wendland covariance function
mu2	one of the smoothness parameters associated with the Generalized Wendland covariance function
family	covariance function family, the options are c("matern", "gw", "cs", "spher", "pexp", "gaussian").
range	Minimum and maximum distance to be considered. The default is range = c(1e-04, 1000).
cut	desired spatial correlation at a distance d, the default is cut = .05.

Value

a numeric value indicating the range parameter such that the spatial correlation between two points at distance d is cut.

fit_spm

Fitting an underlying continuous process to areal data

Description

This function fits a spatial process model to areal data by estimating the parameters of an underlying continuous process. It leverages the [stats::optim](#) function for Maximum Likelihood estimation, providing flexibility in optimization algorithms and control parameters.

Usage

```
fit_spm(x, ...)

## S3 method for class 'spm'
fit_spm(
  x,
  model,
  theta_st,
  nu = NULL,
  kappa = 1,
  mu2 = 1.5,
  apply_exp = FALSE,
  opt_method = "Nelder-Mead",
  control_opt = list(),
  comp_hess = TRUE,
  ...
)

fit_spm2(
  x,
  model,
  nu,
  kappa = 1,
  mu2 = 1.5,
  comp_hess = TRUE,
  phi_min,
  phi_max,
  nphi = 10,
  cores = getOption("mc.cores", 1L)
)
```

Arguments

x An object of type `spm`. The dimension of `theta_st` depends on the number of variables analyzed and whether the input is an `spm` object.

... Additional parameters passed to [stats::optim](#).

model	A character scalar indicating the covariance function family. Options are c("matern", "pexp", "gaussian", "spherical", "gw").
theta_st	A named numeric vector containing initial parameter values.
nu	A numeric value specifying the ν parameter for the Matern or Powered Exponential covariance functions. If model is "matern" and nu is not provided, it defaults to 0.5. If model is "pexp" and nu is not provided, it defaults to 1. In both cases, this results in the exponential covariance function.
kappa	$\kappa \in \{0, \dots, 3\}$ parameter for the GW covariance function.
mu2	The smoothness parameter μ for the GW function.
apply_exp	A logical scalar indicating whether to exponentiate non-negative parameters.
opt_method	A character scalar specifying the optimization algorithm (see stats::optim).
control_opt	A named list of control arguments for the optimization algorithm (see stats::optim).
comp_hess	A logical indicating whether to compute the Hessian matrix.
phi_min	A numeric scalar representing the minimum <i>phi</i> value for grid search.
phi_max	A numeric scalar representing the maximum <i>phi</i> value for grid search.
nphi	A numeric scalar indicating the number of <i>phi</i> values for grid search.
cores	An integer scalar indicating the number of cores to use. Defaults to <code>getOption("mc.cores")</code> . No effect on Windows.

Details

The function utilizes optimization algorithms from [stats::optim](#) to determine Maximum Likelihood estimators (MLEs) and their standard errors for a model adapted for areal data. Users can customize the optimization process by providing control parameters through the `control_opt` argument (a named list, see [stats::optim](#) for details), specifying lower and upper bounds for parameters, and selecting the desired optimization algorithm via `opt_method` (must be available in [stats::optim](#)).

Additionally, the function supports various covariance functions, including Matern, Exponential, Powered Exponential, Gaussian, and Spherical. The `apply_exp` argument, a logical scalar, allows for exponentiation of non-negative parameters, enabling optimization over the entire real line.

The underlying model assumes a point-level process:

$$Y(\mathbf{s}) = \mu + S(\mathbf{s})$$

where $S \sim GP(0, \sigma^2 C(\|\mathbf{s} - \mathbf{s}_2\|; \theta))$. The observed areal data is then assumed to behave as the average of the process over each area:

$$Y(B) = |B|^{-1} \int_B Y(\mathbf{s}) \, d\mathbf{s}$$

Value

An `spm_fit` object containing the estimated model parameters.

Examples

```

data(liv_lsoa) ## loading the LSOA data

msoa_spm <- sf_to_spm(sf_obj = liv_msoa, n_pts = 500,
                    type = "regular", by_polygon = FALSE,
                    poly_ids = "msoa11cd",
                    var_ids = "leb_est")

## fitting model
theta_st_msoa <- c("phi" = 1) # initial value for the range parameter

fit_msoa <-
  fit_spm(x = msoa_spm,
         theta_st = theta_st_msoa,
         model = "matern",
         nu = .5,
         apply_exp = TRUE,
         opt_method = "L-BFGS-B",
         control = list(maxit = 500))

AIC(fit_msoa)

summary_spm_fit(fit_msoa, sig = .05)

```

goodness_of_fit	<i>Akaike's (and Bayesian) An Information Criterion for spm_fit objects.</i>
-----------------	--

Description

Akaike's (and Bayesian) An Information Criterion for `spm_fit` objects.

Usage

```

## S3 method for class 'spm_fit'
AIC(object, ..., k = 2)

## S3 method for class 'spm_fit'
BIC(object, ...)

```

Arguments

object	a <code>spm_fit</code> object.
...	optionally more fitted model objects.
k	numeric, the <i>penalty</i> per parameter to be used; the default 'k = 2' is the classical AIC. (for compatibility with <code>stats::AIC</code> .)

Value

a numeric scalar corresponding to the goodness of fit measure.

liv_lsoa	<i>Liverpool Lower Super Output Area.</i>
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Description

A dataset containing containing the LSOA's for Liverpool along with estimates for Index of Multiple Deprivation. Data taken from [Johnson et al. 2020](#)

Usage

liv_lsoa

Format

A sf data frame with 298 rows and 6 variables:

lsoa11cd LSOA code

lsoa11nm LSOA name

male Male population

female Female population

imdscore Index of Multiple Deprivation

area LMSOA area, in km^2

Details

The data was projected to EPSG 27700 and units changed to km

Source

<https://ij-healthgeographics.biomedcentral.com/articles/10.1186/s12942-020-00200-w>

liv_msoa

Liverpool Middle Super Output Area.

Description

A dataset containing containing the MSOA's for Liverpool along with estimates for Life Expectancy at Birth. Data taken from [Johnson et al. 2020](#)

Usage

liv_msoa

Format

A sf data frame with 61 rows and 4 variables:

msoa11cd MSOA code

msoa11cd MSOA name

lev_est Estimated life expectancy at birth, in years

area MSOA area, in km^2

Details

The data was projected to EPSG 27700 and units changed to km

Source

<https://ij-healthgeographics.biomedcentral.com/articles/10.1186/s12942-020-00200-w>

nl_ct

Nova Lima census tracts

Description

A dataset containing containing the census tracts for the city of Nova Lima in Minas Gerais - Brazil.

Usage

nl_ct

Format

A sf data frame with 113 rows and 14 variables:

cd_setor unique identifier
hh_density average household density
var_hhd variance of the household density
avg_income average income per household
var_income variance of the income per household
pop population in the census tract
avg_age average age of the inhabitants in the census tract
var_age variance of the variable age in the census tract
prop_women proportion of women
prop_elder proportion of people with 55 years of age or older
illit_rate illiteracy rate
prop_white proportion of self-declared white people
prop_black proportion of self-declared black people
prop_native proportion of self-declared native people

Details

The data is project using the SIRGAS 2000.

predict_spm	<i>Prediction over the same or a different set of regions (or points).</i>
-------------	--

Description

Realizes predictions that can be useful when researchers are interested in predict a variable observed in one political division of a city (or state) on another division of the same region.

Usage

```
predict_spm(x, ...)

## S3 method for class 'spm_fit'
predict_spm(x, .aggregate = TRUE, ...)

## S3 method for class 'sf'
predict_spm(x, spm_obj, n_pts, type, outer_poly = NULL, id_var, ...)
```

Arguments

<code>x</code>	a sf object such that its geometries are either points or polygons.
<code>...</code>	additional parameters
<code>.aggregate</code>	logical. Should the predictions be aggregated? In case the input is only a "fit" object, the aggregation is made over the polygons on which the original data was observed. In case the input <code>x</code> is composed by sf POLYGONS, the aggregation is made over this new partition of the study region.
<code>spm_obj</code>	an object of either class <code>spm_fit</code> or <code>mspm_fit</code>
<code>n_pts</code>	a numeric scalar standing for number of points to form a grid over the whole region to make the predictions
<code>type</code>	character type of grid to be generated. See <code>st_sample</code> in the package <code>sf</code> .
<code>outer_poly</code>	(object) sf geometry storing the "outer map" we want to compute the predictions in.
<code>id_var</code>	if <code>x</code> is a set of POLYGONS (areal data) instead of a set of points, the <code>id_var</code> is the name (or index) of the unique identifier associated to each polygon.

Value

a list of size 4 belonging to the class `spm_pred`. This list contains the predicted values and the mean and covariance matrix associated with the conditional distribution used to compute the predictions.

Examples

```
data(liv_lsoa) ## loading the LSOA data
data(liv_msoa) ## loading the MSOA data

msoa_spm <- sf_to_spm(sf_obj = liv_msoa, n_pts = 500,
                    type = "regular", by_polygon = FALSE,
                    poly_ids = "msoa11cd",
                    var_ids = "leb_est")

## fitting model
theta_st_msoa <- c("phi" = 1) # initial value for the range parameter

fit_msoa <-
  fit_spm(x = msoa_spm,
         theta_st = theta_st_msoa,
         model = "matern",
         nu = .5,
         apply_exp = TRUE,
         opt_method = "L-BFGS-B",
         control = list(maxit = 500))

pred_lsoa <- predict_spm(x = liv_lsoa, spm_obj = fit_msoa, id_var = "lsoa11cd")
```

sev_pexp	<i>Calculate Smallest Eigenvalue for Power Exponential Correlation Matrices</i>
----------	---

Description

This function computes the smallest eigenvalue of a correlation matrix derived from the power exponential correlation function. It evaluates this across a grid of values for the power parameter (ν) and the practical range parameter (ρ), based on a provided distance matrix.

Usage

```
sev_pexp(range_nu, range_rho, grid_len = 50, dmat)
```

Arguments

range_nu	A numeric vector of length 2, specifying the minimum and maximum values for the power parameter ν . ν typically ranges between 0 and 2 (e.g., $\nu = 1$ for exponential, $\nu = 2$ for Gaussian).
range_rho	A numeric vector of length 2, specifying the minimum and maximum values for the practical range parameter ρ . ρ must be positive.
grid_len	An integer specifying the number of points to create for both ν and ρ sequences. The total number of grid combinations will be grid_len^2 . Default is 50.
dmat	A numeric matrix representing the distance matrix between locations. The distances should be non-negative.

Details

The practical range ρ is defined here as the distance at which the correlation is 0.1. The internal scale parameter ϕ is calculated as $\phi = \rho / (\log(10)^{1/\nu})$. The power exponential correlation function is assumed to be of the form $C(h) = \exp(-(h/\phi)^\nu)$, where h is distance. The function `smile:::pexp_cov` is used internally to compute the covariance/correlation matrix with a sill of 1.

The function first creates a grid of ν and ρ parameters. For each pair of (ρ, ν) in the grid: 1. It calculates the scale parameter ϕ for the power exponential correlation function, where $\phi = \rho / (\log(10)^{1/\nu})$. This definition implies that the correlation is 0.1 at the distance ρ . 2. It computes the power exponential correlation matrix using `smile:::pexp_cov(dists = dmat, sill = 1, range = phi, smooth = nu)`. Note the use of an internal function from the `smile` package. 3. It calculates the eigenvalues of this correlation matrix. 4. The minimum eigenvalue is extracted. The final output is a tibble containing all parameter combinations and their corresponding minimum eigenvalues.

Value

A **tibble** with three columns:

rho	The practical range parameter value.
nu	The power parameter value.
lambda	The smallest eigenvalue of the power exponential correlation matrix corresponding to the rho and nu pair.

sf_to_spm	<i>single sf to spm</i>
-----------	-------------------------

Description

Transforming a sf into a spm object (Internal use)

Usage

```
single_sf_to_spm(
  sf_obj,
  n_pts,
  type = "regular",
  by_polygon = FALSE,
  poly_ids = NULL,
  var_ids = NULL
)
```

```
sf_to_spm(
  sf_obj,
  n_pts,
  type = "regular",
  by_polygon = FALSE,
  poly_ids = NULL,
  var_ids = NULL
)
```

Arguments

sf_obj	a sf object s.t. its geometries are polygons.
n_pts	a numeric scalar representing the number of points to create a grid in the study region on which the polygons in sf_obj is observed. Alternatively, it can be a vector of the same length as nrow(sf_obj). In this case, it generates the given number of points for each polygon in sf_obj.
type	a character indicating the type of grid to be generated. The options are c("random", "regular", "hexagonal"). For more details, see st_sample in the sf package.

by_polygon	a logical indicating whether we should generate n_pts by polygon or for the n_pts for the whole study region.
poly_ids	a character vector informing the name of the variable in sf_obj that represents the polygons unique identifiers. In case this is not informed, we assume the id of the polygons are given by their row numbers.
var_ids	a scalar or vector of type character indicating the (numerical) variables that are going to be analyzed.

Value

a named list of size 6 belonging to the class `spm`. This list stores all the objects necessary to fit models using the `fit_spm`.

Examples

```
data(liv_lsoa) # loading the LSOA data

msoa_spm <- sf_to_spm(sf_obj = liv_msoa, n_pts = 1000,
                    type = "regular", by_polygon = FALSE,
                    poly_ids = "msoa11cd",
                    var_ids = "leb_est")
```

smile	<i>smile: Spatial MIsaLignment Estimation</i>
-------	---

Description

smile: Spatial MIsaLignment Estimation

summary_spm_fit	<i>Summarizing spm_fit</i>
-----------------	----------------------------

Description

Provides a `data.frame` with point estimates and confidence intervals for the parameters of the model fitted using the `spm_fit` function.

Usage

```
summary_spm_fit(x, sig = 0.05)
```

Arguments

x	a <code>spm_fit</code> object.
sig	a real number between 0 and 1 indicating significance level to be used to compute the confidence intervals for the parameter estimates.

Value

a `data.frame` summarizing the parameters estimated by the `fit_spm` function.

vdl	<i>Voronoi Data Linkage</i>
-----	-----------------------------

Description

Reminder, have to create an example. This will be exported after we submit the paper for publication.

Usage

```
vdl(coords_sf, areal_sf, vars, buff)
```

Arguments

<code>coords_sf</code>	sf POINT target dataset.
<code>areal_sf</code>	sf POLYGON source dataset.
<code>vars</code>	a character representing the variables (observed at the source - polygon) to be estimated at the target data.
<code>buff</code>	scalar numeric. Mostly for internal use.

Value

a sf object for the `coords_sf` spatial data set.

vdl_var	<i>Voronoi Data Linkage - Single variable and variance</i>
---------	--

Description

Reminder, have to create an example. This will be exported after we submit the paper for publication.

Usage

```
vdl_var(coords_sf, areal_sf, res_var, variance, var_method = "CS", buff)
```

Arguments

coords_sf	sf POINT target dataset.
areal_sf	sf POLYGON source dataset.
res_var	a character - the name of the variable in the areal_sf to be estimated in the coords_sf.
variance	a character - the name of the variable variance in the areal_sf to be estimated in the coords_sf.
var_method	a character representing the method to approximate the variance of the AI estimates. Possible values are "CS" (Cauchy-Schwartz) or "MI" (Moran's I).
buff	scalar numeric. Mostly for internal use.

Value

a sf object, containing the id_coords variable and the list_vars for the coords_sf spatial data set.

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