

# Package ‘sdsfun’

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**Title** Spatial Data Science Complementary Features

**Version** 0.8.0

**Description** Wrapping and supplementing commonly used functions in the R ecosystem related to spatial data science,  
while serving as a basis for other packages maintained by Wenbo Lv.

**License** GPL-3

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## Contents

check_tbl_na . . . . .	2
cor_test . . . . .	3
discretize_vector . . . . .	4
dummy_tbl . . . . .	5

dummy_vec . . . . .	5
formula_varname . . . . .	6
fuzzyoverlay . . . . .	6
generate_subsets . . . . .	7
geodetector_q . . . . .	8
hclustgeo_disc . . . . .	8
inverse_distance_swm . . . . .	9
loess_optnum . . . . .	10
moran_test . . . . .	11
normalize_vector . . . . .	12
rm_lineartrend . . . . .	12
sf_coordinates . . . . .	13
sf_distance_matrix . . . . .	14
sf_geometry_name . . . . .	14
sf_geometry_type . . . . .	15
sf_gk_proj_cgcs2000 . . . . .	15
sf_utm_proj_wgs84 . . . . .	16
sf_voronoi_diagram . . . . .	17
spade_psd . . . . .	17
spdep_contiguity_swm . . . . .	18
spdep_distance_swm . . . . .	19
spdep_lmtest . . . . .	21
spdep_nb . . . . .	21
spdep_skater . . . . .	22
spvar . . . . .	23
ssh_test . . . . .	24
standardize_vector . . . . .	24
tbl_all2int . . . . .	25
tbl_xyz2mat . . . . .	25

**Index****27**


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check_tbl_na	<i>check for NA values in a tibble</i>
--------------	--

---

**Description**

check for NA values in a tibble

**Usage**

```
check_tbl_na(tbl)
```

**Arguments**

tbl	A tibble
-----	----------

**Value**

A logical value.

**Examples**

```
demotbl = tibble::tibble(x = c(1,2,3,NA,1),
                         y = c(NA,NA,1:3),
                         z = 1:5)
demotbl
check_tbl_na(demotbl)
```

---

cor_test	(partial) correlation test
----------	----------------------------

---

**Description**

(partial) correlation test

**Usage**

```
cor_test(x, y, z = NULL, level = 0.05)
```

**Arguments**

- x A numeric vector representing the first variable.
- y A numeric vector representing the second variable.
- z An optional numeric vector or matrix of control variables. If provided, partial correlation is computed.
- level (optional) Significance level. Default is 0.05.

**Value**

A numeric vector

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
cor_test(gzma$PS_Score, gzma$EL_Score)
```

discretize_vector	<i>discretization</i>
-------------------	-----------------------

## Description

`discretization`

## Usage

```
discretize_vector(
  x,
  n,
  method = "natural",
  breakpoint = NULL,
  sampleprob = 0.15,
  thr = 0.4,
  seed = 123456789
)
```

## Arguments

<code>x</code>	A continuous numeric vector.
<code>n</code>	(optional) The number of discretized classes.
<code>method</code>	(optional) The method of discretization, default is <code>natural</code> .
<code>breakpoint</code>	(optional) Break points for manually splitting data. When <code>method</code> is <code>manual</code> , <code>breakpoint</code> is required.
<code>sampleprob</code>	(optional) When the data size exceeds 3000, perform sampling for discretization, applicable only to natural breaks. Default is 0.15.
<code>thr</code>	(optional) Threshold for controlling iteration, applicable only to headtails breaks. Default is 0.15.
<code>seed</code>	(optional) Random seed number, default is 123456789.

## Value

A discretized integer vector

## Examples

```
xvar = c(22361, 9573, 4836, 5309, 10384, 4359, 11016, 4414, 3327, 3408,
       17816, 6909, 6936, 7990, 3758, 3569, 21965, 3605, 2181, 1892,
       2459, 2934, 6399, 8578, 8537, 4840, 12132, 3734, 4372, 9073,
       7508, 5203)
discretize_vector(xvar, n = 5, method = 'natural')
```

---

dummy_tbl	<i>transforming a category tibble into the corresponding dummy variable tibble</i>
-----------	--

---

**Description**

transforming a category tibble into the corresponding dummy variable tibble

**Usage**

```
dummy_tbl(tbl)
```

**Arguments**

tbl            A tibble or data.frame.

**Value**

A tibble

**Examples**

```
a = tibble::tibble(x = 1:3,y = 4:6)
dummy_tbl(a)
```

---

dummy_vec	<i>transforming a categorical variable into dummy variables</i>
-----------	---

---

**Description**

transforming a categorical variable into dummy variables

**Usage**

```
dummy_vec(x)
```

**Arguments**

x            An integer vector or can be converted into an integer vector.

**Value**

A matrix.

**Examples**

```
dummy_vec(c(1,1,3,2,4,6))
```

---

<code>formula_varname</code>	<i>get variable names in a formula and data</i>
------------------------------	---

---

### Description

get variable names in a formula and data

### Usage

```
formula_varname(formula, data)
```

### Arguments

<code>formula</code>	A formula.
<code>data</code>	A <code>data.frame</code> , <code>tibble</code> or <code>sf</code> object of observation data.

### Value

A list.

`yname` Independent variable name

`xname` Dependent variable names

### Examples

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
formula_varname(PS_Score ~ EL_Score + OH_Score, gzma)
formula_varname(PS_Score ~ ., gzma)
```

---

<code>fuzzyoverlay</code>	<i>spatial fuzzy overlay</i>
---------------------------	------------------------------

---

### Description

spatial fuzzy overlay

### Usage

```
fuzzyoverlay(formula, data, method = "and")
```

### Arguments

<code>formula</code>	A formula of spatial fuzzy overlay.
<code>data</code>	A <code>data.frame</code> or <code>tibble</code> of discretized data.
<code>method</code>	(optional) Overlay methods. When <code>method</code> is <code>and</code> , use <code>min</code> to do fuzzy overlay; and when <code>method</code> is <code>or</code> , use <code>max</code> to do fuzzy overlay. Default is <code>and</code> .

**Value**

A numeric vector.

**Note**

Independent variables in the data provided to `fuzzyoverlay()` must be discretized variables, and dependent variable are continuous variable.

**Examples**

```
set.seed(42)
sim = tibble::tibble(y = stats::runif(7,0,10),
                     x1 = c(1,rep(2,3),rep(3,3)),
                     x2 = c(rep(1,2),rep(2,2),rep(3,3)))
fo1 = fuzzyoverlay(y~x1+x2,data = sim, method = 'and')
fo1
fo2 = fuzzyoverlay(y~x1+x2,data = sim, method = 'or')
fo2
```

generate_subsets	<i>generate subsets of a set</i>
------------------	----------------------------------

**Description**

generate subsets of a set

**Usage**

```
generate_subsets(set, empty = TRUE, self = TRUE)
```

**Arguments**

<code>set</code>	A vector.
<code>empty</code>	(optional) When <code>empty</code> is TRUE, the generated subset includes the empty set, otherwise the empty set is removed. Default is TRUE.
<code>self</code>	(optional) When <code>self</code> is TRUE, the resulting subset includes the set itself, otherwise the set itself is removed. Default is TRUE.

**Value**

A list.

**Examples**

```
generate_subsets(letters[1:3])
generate_subsets(letters[1:3],empty = FALSE)
generate_subsets(letters[1:3],self = FALSE)
generate_subsets(letters[1:3],empty = FALSE,self = FALSE)
```

<code>geodetector_q</code>	<i>only geodetector q-value</i>
----------------------------	---------------------------------

### Description

only geodetector q-value

### Usage

```
geodetector_q(y, hs)
```

### Arguments

<code>y</code>	Dependent variable
<code>hs</code>	Independent variable

### Value

A numeric value

### Examples

```
geodetector_q(y = 1:7, hs = c('x',rep('y',3),rep('z',3)))
```

<code>hclustgeo_disc</code>	<i>hierarchical clustering with spatial soft constraints</i>
-----------------------------	--

### Description

hierarchical clustering with spatial soft constraints

### Usage

```
hclustgeo_disc(
  data,
  n,
  alpha = 0.5,
  D1 = NULL,
  hclustm = "ward.D2",
  scale = TRUE,
  wt = NULL,
  ...
)
```

## Arguments

data	An <code>sf</code> object, <code>tibble</code> , <code>data.frame</code> , <code>matrix</code> or <code>vector</code> of observations data.
n	The number of hierarchical clustering classes, which can be a numeric value or vector.
alpha	(optional) A positive value between 0 and 1. This mixing parameter gives the relative importance of "feature" space and "constraint" space. Default is 0.5.
D1	(optional) A <code>matrix</code> with other dissimilarities between the same observations data. if <code>data</code> is an <code>sf</code> object and <code>alpha</code> is not 0, the <code>D1</code> will be generated by <code>sdsfun::sf_distance_matrix()</code> , others will use a <code>matrix</code> with all elements equal to 0.
hclustm	(optional) The agglomeration method to be used, default is <code>ward.D2</code> . For more details, please see <code>stats::hclust()</code> .
scale	(optional) Whether to scaled the dissimilarities matrix, default is TRUE.
wt	(optional) Vector with the weights of the observations. By default, <code>wt</code> is NULL.
...	(optional) Other arguments passed to <code>stats::dist()</code> .

## Value

The grouped membership: a `vector` if `n` is a scalar, a `matrix` (columns correspond to elements of `n`) if not.

## Note

This is a C++ enhanced implementation of the `hclustgeo` function in `ClustGeo` package.

## Examples

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
gzma$group = hclustgeo_disc(gzma, 5, alpha = 0.75)
plot(gzma["group"])
```

`inverse_distance_swm` *construct inverse distance weight*

## Description

Function for constructing inverse distance weight.

## Usage

```
inverse_distance_swm(sfj, power = 1, bandwidth = NULL)
```

**Arguments**

- sfj** Vector object that can be converted to sf by `sf::st_as_sf()`.  
**power** (optional) Default is 1. Set to 2 for gravity weights.  
**bandwidth** (optional) When the distance is bigger than bandwidth, the corresponding part of the weight matrix is set to 0. Default is NULL, which means not use the bandwidth.

**Details**

The inverse distance weight formula is  $w_{ij} = 1/d_{ij}^\alpha$

**Value**

A inverse distance weight matrices with class of `matrix`.

**Examples**

```
library(sf)
pts = read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))
wt = inverse_distance_swm(pts)
wt[1:5,1:5]
```

**loess\_optnum**

*determine optimal spatial data discretization for individual variables*

**Description**

Function for determining optimal spatial data discretization for individual variables based on locally estimated scatterplot smoothing (LOESS) model.

**Usage**

```
loess_optnum(qvec, discnumvec, increase_rate = 0.05)
```

**Arguments**

- qvec** A numeric vector of q statistics.  
**discnumvec** A numeric vector of break numbers corresponding to qvec.  
**increase\_rate** (optional) The critical increase rate of the number of discretization. Default is 0.05.

**Value**

A two element numeric vector.

**discnum** optimal number of spatial data discretization

**increase\_rate** the critical increase rate of the number of discretization

**Note**

When `increase_rate` is not satisfied by the calculation, the discrete number corresponding to the highest `q_statistic` is selected as a return.

Note that `sdsfun` sorts `discnumvec` from smallest to largest and keeps `qvec` in one-to-one correspondence with `discnumvec`.

**Examples**

```
qv = c(0.26045642, 0.64120405, 0.43938704, 0.95165535, 0.46347836,
      0.25385338, 0.78778726, 0.95938330, 0.83247885, 0.09285196)
loess_optnum(qv, 3:12)
```

moran\_test

*global spatial autocorrelation test***Description**

global spatial autocorrelation test

**Usage**

```
moran_test(sfj, wt = NULL, alternative = "greater", symmetrize = FALSE)
```

**Arguments**

<code>sfj</code>	An <code>sf</code> object or can be converted to <code>sf</code> by <code>sf::st_as_sf()</code> .
<code>wt</code>	(optional) Spatial weight matrix. Must be a <code>matrix</code> class. If <code>wt</code> is not provided, <code>sdsfun</code> will use a first-order queen adjacency binary matrix.
<code>alternative</code>	(optional) Specification of alternative hypothesis as <code>greater</code> (default), <code>lower</code> , or <code>two.sided</code> .
<code>symmetrize</code>	(optional) Whether or not to symmetrize the asymmetrical spatial weight matrix <code>wt</code> by: $1/2 * (\mathbf{wt} + \mathbf{wt}')$ . Default is <code>FALSE</code> .

**Value**

A list utilizing a `result tibble` to store the following information for each variable:

`MoranI` observed value of the Moran coefficient

`EI` expected value of Moran's I

`VarI` variance of Moran's I (under normality)

`ZI` standardized Moran coefficient

`PI` *p*-value of the test statistic

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
moran_test(gzma)
```

normalize_vector	<i>normalization</i>
------------------	----------------------

**Description**

normalization

**Usage**

```
normalize_vector(x, to_left = 0, to_right = 1)
```

**Arguments**

- x                A continuous numeric vector.
- to\_left         (optional) Specified minimum. Default is 0.
- to\_right        (optional) Specified maximum. Default is 1.

**Value**

A continuous vector which has normalized.

**Examples**

```
normalize_vector(c(-5,1,5,0.01,0.99))
```

rm_lineartrend	<i>remove variable linear trend based on covariate</i>
----------------	--

**Description**

remove variable linear trend based on covariate

**Usage**

```
rm_lineartrend(formula, data, method = c("cpp", "r"))
```

**Arguments**

formula	A formula.
data	The observation data.
method	(optional) The method for using, which can be chosen as either <code>cpp</code> or <code>r</code> . Default is <code>cpp</code> .

**Value**

A numeric vector.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))  
rm_lineartrend(PS_Score ~ ., gzma)  
rm_lineartrend(PS_Score ~ ., gzma, method = "r")
```

---

sf_coordinates	<i>extract locations</i>
----------------	--------------------------

---

**Description**

Extract locations of sf objects.

**Usage**

```
sf_coordinates(sfj)
```

**Arguments**

sfj	An sf object or can be converted to sf by <code>sf::st_as_sf()</code> .
-----	---

**Value**

A matrix.

**Examples**

```
pts = sf::read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))  
sf_coordinates(pts)
```

`sf_distance_matrix`     *generates distance matrix*

### Description

Generates distance matrix for sf object

### Usage

```
sf_distance_matrix(sfj)
```

### Arguments

`sfj`     An sf object or can be converted to sf by `sf::st_as_sf()`.

### Value

A matrix.

### Examples

```
pts = sf::read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))
pts_distm = sf_distance_matrix(pts)
pts_distm[1:5,1:5]
```

`sf_geometry_name`     *sf object geometry column name*

### Description

Get the geometry column name of an sf object

### Usage

```
sf_geometry_name(sfj)
```

### Arguments

`sfj`     An sf object.

### Value

A character.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
sf_geometry_name(gzma)
```

`sf_geometry_type`      *sf object geometry type*

**Description**

Get the geometry type of an `sf` object

**Usage**

```
sf_geometry_type(sfj)
```

**Arguments**

`sfj`      An `sf` object.

**Value**

A lowercase character vector

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
sf_geometry_type(gzma)
```

`sf_gk_proj_cgcs2000`      *generates cgcs2000 Gauss-Kruger projection epsg coding character*

**Description**

Generates a Gauss-Kruger projection epsg coding character corresponding to an `sfj` object under the CGCS2000 spatial reference.

**Usage**

```
sf_gk_proj_cgcs2000(sfj, degree = 6L)
```

**Arguments**

<code>sfj</code>	An <code>sf</code> object or can be converted to <code>sf</code> by <code>sf::st_as_sf()</code> .
<code>degree</code>	(optional) 3-degree or 6-degree zonal projection, default is 6L.

**Value**

A character.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun')) |>
  sf::st_transform(4490)
sf_gk_proj_cgcs2000(gzma, 3)
sf_gk_proj_cgcs2000(gzma, 6)
```

*sf\_utm\_proj\_wgs84*      generates wgs84 utm projection epsg coding character

**Description**

Generates a utm projection epsg coding character corresponding to an *sfj* object under the WGS84 spatial reference.

**Usage**

```
sf_utm_proj_wgs84(sfj)
```

**Arguments**

sfj	An <i>sf</i> object or can be converted to <i>sf</i> by <i>sf::st_as_sf()</i> .
-----	---

**Value**

A character.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
sf_utm_proj_wgs84(gzma)
```

`sf_voronoi_diagram`     *generates voronoi diagram*

### Description

Generates Voronoi diagram (Thiessen polygons) for sf object

### Usage

```
sf_voronoi_diagram(sfj)
```

### Arguments

sfj	An sf object.
-----	---------------

### Value

An sf object of polygon geometry type or can be converted to this by `sf::st_as_sf()`.

### Note

Only sf objects of (multi-)point type are supported to generate voronoi diagram and the returned result includes only the geometry column.

### Examples

```
pts = sf::read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))
pts_v = sf_voronoi_diagram(pts)

library(ggplot2)
ggplot() +
  geom_sf(data = pts_v, color = 'red',
          fill = 'transparent') +
  geom_sf(data = pts, color = 'blue', size = 1.25) +
  theme_void()
```

`spade_psd`     *only spade power of spatial determinant*

### Description

only spade power of spatial determinant

### Usage

```
spade_psd(y, hs, wt)
```

**Arguments**

y	Dependent variable
hs	Independent variable
wt	Spatial weight matrix

**Value**

A numeric value

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
wt1 = inverse_distance_swm(gzma)
spade_psd(y = gzma$PS_Score,
           hs = discretize_vector(gzma$PS_Score, 5),
           wt = wt1)
```

*spdep\_contiguity\_swm* constructs spatial weight matrices based on contiguity

**Description**

Constructs spatial weight matrices based on contiguity via spdep package.

**Usage**

```
spdep_contiguity_swm(
  sfj,
  queen = TRUE,
  k = NULL,
  order = 1L,
  cumulate = TRUE,
  style = "W",
  zero.policy = TRUE
)
```

**Arguments**

sfc	An sf object or can be converted to sf by sf::st_as_sf().
queen	(optional) if TRUE, using queen contiguity, otherwise rook contiguity. Default is TRUE.
k	(optional) The number of nearest neighbours. Ignore this parameter when not using distance based neighbours to construct spatial weight matrices.
order	(optional) The order of the adjacency object. Default is 1.
cumulate	(optional) Whether to accumulate adjacency objects. Default is TRUE.

style	(optional) style can take values W, B, C, and S. More to see <code>spdep::nb2mat()</code> . Default is W.
zero.policy	(optional) if FALSE stop with error for any empty neighbour sets, if TRUE permit the weights list to be formed with zero-length weights vectors. Default is TRUE.

**Value**

A matrix

**Note**

When k is set to a positive value, using K-Nearest Neighbor Weights.

**Examples**

```
gzma = sf::read_sf(system.file('exdata/gzma.gpkg', package = 'sdsfun'))
wt1 = spdep_contiguity_swm(gzma, k = 6, style = 'B')
wt2 = spdep_contiguity_swm(gzma, queen = TRUE, style = 'B')
wt3 = spdep_contiguity_swm(gzma, queen = FALSE, order = 2, style = 'B')
```

`spdep_distance_swm` constructs spatial weight matrices based on distance

**Description**

Constructs spatial weight matrices based on distance via `spdep` package.

**Usage**

```
spdep_distance_swm(
  sfj,
  kernel = NULL,
  k = NULL,
  bandwidth = NULL,
  power = 1,
  style = "W",
  zero.policy = TRUE
)
```

**Arguments**

<code>sfj</code>	An <code>sf</code> object or can be converted to <code>sf</code> by <code>sf::st_as_sf()</code> .
<code>kernel</code>	(optional) The kernel function, can be one of uniform, triangular,quadratic(epanechnikov),quartic and gaussian. Default is NULL.
<code>k</code>	(optional) The number of nearest neighbours. Default is NULL. Only useful when <code>kernel</code> is provided.

bandwidth	(optional) The bandwidth, default is NULL. When the spatial reference of sf object is the geographical coordinate system, the unit of bandwidth is km. The unit used in the projection coordinate system are consistent with those used in the sf object coordinate system.
power	(optional) Default is 1. Useful when kernel is not provided.
style	(optional) style can take values W, B, C, and S. More to see <code>spdep::nb2mat()</code> . Default is W. For spatial weights based on distance functions, a style of B means using the original value of the calculated distance function.
zero.policy	(optional) if FALSE stop with error for any empty neighbour sets, if TRUE permit the weights list to be formed with zero-length weights vectors. Default is TRUE.

## Details

five different kernel weight functions:

- uniform:  $K(z) = 1/2$ , for  $|z| < 1$
- triangular  $K(z) = 1 - |z|$ , for  $|z| < 1$
- quadratic (epanechnikov)  $K(z) = \frac{3}{4}(1 - z^2)$ , for  $|z| < 1$
- quartic  $K(z) = \frac{15}{16}(1 - z^2)^2$ , for  $|z| < 1$
- gaussian  $K(z) = \frac{1}{\sqrt{2\pi}}e^{-\frac{z^2}{2}}$

For the equation above,  $z = d_{ij}/h_i$  where  $h_i$  is the bandwidth

## Value

A matrix

## Note

When kernel is setting, using distance weight based on kernel function, Otherwise the inverse distance weight will be used.

## Examples

```
pts = sf::read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))
wt1 = spdep_distance_swm(pts, style = 'B')
wt2 = spdep_distance_swm(pts, kernel = 'gaussian')
wt3 = spdep_distance_swm(pts, k = 3, kernel = 'gaussian')
wt4 = spdep_distance_swm(pts, k = 3, kernel = 'gaussian', bandwidth = 10000)
```

---

spdep_lmtest	<i>spatial linear models selection</i>
--------------	--

---

## Description

spatial linear models selection

## Usage

```
spdep_lmtest(formula, data, listw = NULL)
```

## Arguments

- |         |   |
|---------|---|
| formula | A formula for linear regression model.  |
| data    | An sf object of observation data.   |
| listw   | (optional) A listw. See spdep::mat2listw() and spdep::nb2listw() for details. |

## Value

A list

## Examples

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))  
spdep_lmtest(PS_Score ~ ., gzma)
```

---

spdep_nb	<i>construct neighbours list</i>
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---

## Description

construct neighbours list

## Usage

```
spdep_nb(sfj, queen = TRUE, k = NULL, order = 1L, cumulate = TRUE)
```

**Arguments**

<code>sfj</code>	An <code>sf</code> object or can be converted to <code>sf</code> by <code>sf::st_as_sf()</code> .
<code>queen</code>	(optional) if TRUE, using queen contiguity, otherwise rook contiguity. Default is TRUE.
<code>k</code>	(optional) The number of nearest neighbours. Ignore this parameter when not using distance based neighbours.
<code>order</code>	(optional) The order of the adjacency object. Default is 1.
<code>cumulate</code>	(optional) Whether to accumulate adjacency objects. Default is TRUE.

**Value**

A neighbours list with class `nb`

**Note**

When `k` is set to a positive value, using K-Nearest Neighbor

**Examples**

```
pts = sf::read_sf(system.file('extdata/pts.gpkg', package = 'sdsfun'))
nb1 = spdep_nb(pts, k = 6)
nb2 = spdep_nb(pts, queen = TRUE)
nb3 = spdep_nb(pts, queen = FALSE, order = 2)
```

---

`spdep_skater`

*spatial c(k)luster analysis by tree edge removal*

---

**Description**

SKATER forms clusters by spatially partitioning data that has similar values for features of interest.

**Usage**

```
spdep_skater(sfj, k = 6, nb = NULL, ini = 5, ...)
```

**Arguments**

<code>sfj</code>	An <code>sf</code> object of observation data. Please ensure that the attribute columns are included in the SKATER analysis.
<code>k</code>	(optional) The number of clusters. Default is 6.
<code>nb</code>	(optional) A neighbours list with class <code>nb</code> . If the input <code>nb</code> is <code>NULL</code> , it will be constructed automatically using <code>spdep_nb()</code> .
<code>ini</code>	(optional) The initial node in the minimal spanning tree. Default is 5.
<code>...</code>	(optional) Other parameters passed to <code>spdep::skater()</code> .

**Value**

A numeric vector of clusters.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
gzma_c = spdep_skater(gzma, 8)
gzma$group = gzma_c
plot(gzma["group"])
```

spvar

*spatial variance***Description**

spatial variance

**Usage**

```
spvar(x, wt, method = c("cpp", "r"))
```

**Arguments**

- |                     |  |
|---------------------|--|
| <code>x</code>      | A numerical vector .   |
| <code>wt</code>     | The spatial weight matrix.   |
| <code>method</code> | (optional) The method for calculating spatial variance, which can be chosen as either <code>cpp</code> or <code>r</code> . Default is <code>cpp</code> . |

**Details**

The spatial variance formula is  $\Gamma = \frac{\sum_i \sum_{j \neq i} \omega_{ij} \frac{(y_i - y_j)^2}{2}}{\sum_i \sum_{j \neq i} \omega_{ij}}$

**Value**

A numerical value.

**Examples**

```
gzma = sf::read_sf(system.file('extdata/gzma.gpkg', package = 'sdsfun'))
wt1 = inverse_distance_sw(gzma)
spvar(gzma$PS_Score, wt1)
```

<code>ssh_test</code>	<i>spatial stratified heterogeneity test</i>
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### Description

`spatial stratified heterogeneity test`

### Usage

```
ssh_test(y, hs)
```

### Arguments

<code>y</code>	Variable Y, continuous numeric vector.
<code>hs</code>	Spatial stratification or classification of each explanatory variable. <code>factor</code> , <code>character</code> , <code>integer</code> or <code>data.frame</code> , <code>tibble</code> and <code>sf</code> object.

### Value

A `tibble`

### Examples

```
ssh_test(y = 1:7, hs = c('x',rep('y',3),rep('z',3)))
```

<code>standardize_vector</code>	<i>standardization</i>
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### Description

To calculate the Z-score using variance normalization, the formula is as follows:

$$Z = \frac{(x - \text{mean}(x))}{\text{sd}(x)}$$

### Usage

```
standardize_vector(x)
```

### Arguments

<code>x</code>	A numeric vector
----------------	------------------

### Value

A standardized numeric vector

**Examples**

```
standardize_vector(1:10)
```

tbl\_all2int

*convert discrete variables in a tibble to integers***Description**

convert discrete variables in a tibble to integers

**Usage**

```
tbl_all2int(tbl)
```

**Arguments**

tbl            A tibble,data.frame or sf object.

**Value**

A converted tibble,data.frame or sf object.

**Examples**

```
demotbl = tibble::tibble(x = c(1,2,3,3,1),
                         y = letters[1:5],
                         z = c(1L,1L,2L,2L,3L),
                         m = factor(letters[1:5],levels = letters[5:1]))
tbl_all2int(demotbl)
```

tbl\_xyz2mat

*convert xyz tbl to matrix***Description**

convert xyz tbl to matrix

**Usage**

```
tbl_xyz2mat(tbl, x = 1, y = 2, z = 3)
```

**Arguments**

tbl	A <code>tibble</code> , <code>data.frame</code> or <code>sf</code> object.
x	(optional) The x-axis coordinates column number, default is 1.
y	(optional) The y-axis coordinates column number, default is 2.
z	(optional) The z (attribute) coordinates column number, default is 3.

**Value**

A list.

**z\_attrs\_matrix** A matrix with attribute information.

**x\_coords\_matrix** A matrix with the x-axis coordinates.

**y\_coords\_matrix** A matrix with the y-axis coordinates.

**Examples**

```
set.seed(42)
lon = rep(1:3,each = 3)
lat = rep(1:3,times = 3)
zattr = rnorm(9, mean = 10, sd = 1)
demodf = data.frame(x = lon, y = lat, z = zattr)
demodf
tbl_xyz2mat(demodf)
```

# Index

check\_tbl\_na, 2  
cor\_test, 3  
  
discretize\_vector, 4  
dummy\_tbl, 5  
dummy\_vec, 5  
  
formula\_varname, 6  
fuzzyoverlay, 6  
  
generate\_subsets, 7  
geodetector\_q, 8  
  
hclustgeo\_disc, 8  
  
inverse\_distance\_swm, 9  
  
loess\_optnum, 10  
  
moran\_test, 11  
  
normalize\_vector, 12  
  
rm\_lineartrend, 12  
  
sf\_coordinates, 13  
sf\_distance\_matrix, 14  
sf\_geometry\_name, 14  
sf\_geometry\_type, 15  
sf\_gk\_proj\_cgcs2000, 15  
sf\_utm\_proj\_wgs84, 16  
sf\_voronoi\_diagram, 17  
spade\_psd, 17  
spdep\_contiguity\_swm, 18  
spdep\_distance\_swm, 19  
spdep\_lmtest, 21  
spdep\_nb, 21  
spdep\_skater, 22  
spvar, 23  
ssh\_test, 24  
standardize\_vector, 24  
  
tbl\_all2int, 25  
tbl\_xyz2mat, 25