

# Package ‘overlapptest’

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**Title** Test Overlapping of Polygons Against Random Rotation  
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**Description** Tests the observed overlapping polygon area in a collection of polygons against a null model of random rotation, as explained in De la Cruz et al. (2017) <[doi:10.13140/RG.2.2.12825.72801](https://doi.org/10.13140/RG.2.2.12825.72801)>.  
**License** GPL (>= 2)  
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Astragalus

*Astragalus and Sesleria Plants*


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### Description

Oulines of the individuals of *Astragalus sempervirens* and *Sesleria albicans* present in a 2 x 2 m plot in the Pyrenean mountains.

### Usage

```
data("Astragalus")
data("Sesleria")
```

### Format

Each object is a multiple-polygon window with the format `owin` of **spatstat**.

### Examples

```
data(Astragalus)
Astragalus
plot(Astragalus)
# total area covered by Astragalus
area.owin(Astragalus)
# number of individual polygons
length(Astragalus$bdry)
# area of each individual
sapply(Astragalus$bdry, function(x) area.owin(owin(poly=x)))
```

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centroidiam

*Compute Centroids and Diameters*


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### Description

Computes the centroid and diameter of each individual polygon in a multi-polygon `owin` object.

### Usage

```
centroidiam(ventana1)
```

### Arguments

ventana1      A multiple-polygon window with the format `owin` of **spatstat**

### Details

Iteratively applies the functions `centroid.owin` and `diameter` of **spatstat** to each polygon in the multipolygon `owin` and computes its centroid and its diameter.

**Value**

diams	Vector of diameters
centroids	Matrix with the coordinates of the centroids

**Author(s)**

Marcelino de la Cruz Rot

**Examples**

```
data(Astragalus)
X<-centroidiam(Astragalus)
X$centroids
X$diams
```

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check.ventana	<i>Checks for Anticlockwise Vertices</i>
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**Description**

Checks that the vertices of polygons in a multi-polygon [owin](#) object are listed anticlockwise and, if some are not, tries to correct them.

**Usage**

```
check.ventana(ventana)
```

**Arguments**

ventana      A multiple-polygon window with the format [owin](#) of **spatstat**.

**Details**

This functions should be employed after importing shapefiles into [owin](#) objects (see vignette). Vertices of the individual polygons in the multiple-polygon [owin](#) objects should be listed anticlockwise to avoid errors in the computations of area overlapp (clockwise listed polygons represent "holes" in **spatstat**). This functions checks this and, in case that the vertices of some polygons are listed clockwise, tries to revert their order.

**Value**

A multiple-polygon window with the format [owin](#) of **spatstat** with the vertices of all polygons listed anticlockwise. The order number of the corrected polygons are included in the attribute "malos1". If there has been any polygon whose vertices have not been corrected, their orden number are included in the attribute "malos2": these polygons should be corrected manually.

**Author(s)**

Marcelino de la Cruz Rot

**Examples**

```
data(Astragalus)
# For illustrative purposes, make the vertices of some individual polygon to be listed clockwise
Astragalus.malo<-Astragalus
Astragalus.malo$bdry[[14]]<-lapply(Astragalus.malo$bdry[[14]], rev)
# check and correct
Astragalus.corrected<-(check.ventana(Astragalus.malo))
attributes(Astragalus.corrected)
```

---

pval

*P Value for a Monte Carlo Test of Polygon Overlapping*

---

**Description**

Computes the p-value and sign of deviation for the hypothesis that the first value in a vector is larger or smaller (i.e., a two-sided test) than the expected value represented by all the other values in the vector, or alternatively, computes a one-sided test.

**Usage**

```
pval(x, alternative=c("two.sided", "less", "greater"))
```

**Arguments**

x	A vector, usually with some observed statistic in the first position (the observed overlap) followed by a sequence of the same statistic computed for several realizations of a null model against which we test our hypothesis (i.e., a sequence of simulated, i.e., rotated, overlaps).
alternative	a character string specifying the alternative hypothesis, must be one of "two.sided" (default), "greater" or "less". You can specify just the initial letter.

**Value**

Vector of length one whose absolute value represents the p-value and whose sign indicates whether the first value in x is larger (positive) or smaller (negative) than the expected value.

**Author(s)**

Marcelino de la Cruz

**Examples**

```
pval(c(0,1:99))
pval(c(100,1:99))
pval(c(100,1:199))
pval(c(200,1:199))
pval(c(0,1:199))
```

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**rotawin***Rotate Individual Polygons*

---

**Description**

Randomly rotates individual polygons around their centroids.

**Usage**

```
rotawin(ventana)
```

**Arguments**

ventana            A multiple-polygon window with the format [owin](#) of **spatstat**.

**Details**

rotawin applies an independent random rotation to each of the polygons in a multiple-polygon [owin](#) object.

**Value**

rotawin returns the original [owin](#) object with the individual polygons randomly rotated.

**Author(s)**

Marcelino de la Cruz Rot

**Examples**

```
data(Astragalus)
plot(Astragalus)
plot(rotawin(Astragalus), add=TRUE, border=2)
```

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test.intersection	<i>Test Overlapping of Polygons Against Random Rotation</i>
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### Description

These functions test the overlapping surface area of a collection of polygons against a null model of random rotation. `test.auto.intersection` test for overlapping between polygons of the same type (i.e., the same species) whereas `test.intersection` test for overlapping between polygons of different types (i.e., between different species). `test.auto.intersection.p` and `test.intersection.p` functions are parallelized versions which might significantly reduce computing times.

### Usage

```
test.auto.intersection(ventana1, nsim = 199, win = NULL,
  prop=1, centroides1= NULL,   diametros1 =NULL)
test.intersection(ventana1, ventana2, nsim, prop = 1, win = NULL,
  centroides1 = NULL, diametros1 = NULL, centroides2 = NULL,
  diametros2 = NULL)
test.auto.intersection.p(ventana1, nsim = 199, win = NULL,
  prop=1, centroides1= NULL,   diametros1 =NULL, ncl=2)
test.intersection.p(ventana1, ventana2, nsim, prop = 1, win = NULL,
  centroides1 = NULL, diametros1 = NULL, centroides2 = NULL,
  diametros2 = NULL, ncl=2)
```

### Arguments

ventana1	A multiple-polygon window with the format <code>owin</code> of <b>spatstat</b> . The "accessory" species in the case of <code>test.intersection</code> .
ventana2	A multiple-polygon window with the format <code>owin</code> of <b>spatstat</b> . The species whose polygons will be rotated (i.e., the "focal" species) in the case of <code>test.intersection</code> .
nsim	Number of simulations for the Monte Carlo test.
prop	Proportion of the diameter of each polygon which will be employed to select potential close neighbours. See details.
win	Observation window, employed to control for edge effects. An object with the format <code>owin</code> of <b>spatstat</b> . If not provided, it will be estimated by the $x$ and $y$ ranges from <code>ventana</code> .
centroides1	Matrix or data.frame with the coordinates of the centroids of the polygons of <code>ventana1</code> .
diametros1	Vector with the diameters of the polygons of <code>ventana1</code> .
centroides2	Matrix or data.frame with the coordinates of the centroids of the polygons of <code>ventana2</code> .
diametros2	Vector with the diameters of the polygons of <code>ventana2</code> .
ncl	Number of clusters for the parallel implementation.

## Details

The summary statistic employed in the test is the overall sum of the overlapping areas of intersecting polygons. To reduce the computing burden, area intersections are only computed for polygon pairs for which the distance between their centroids is smaller than the sum of their largest diameters (weighted by the argument `prop`). To avoid edge effects, the test only considers polygons whose centroids are separated from the border of the observation window by a distance larger than their largest diameter. In the pairwise tests, the polygons of `ventana1` are kept fixed in their original positions (i.e., the accessory species) and the polygons of `ventana2` (i.e., the focal species) are rotated.

## Value

Both `test.auto.intersection` and `test.intersection` return a vector of length `nsim+1`, with the sum of observed overlapping areas in the first position and subsequently with the sum of overlapping areas in each the simulated (i.e., randomly rotated) realizations of the null model.

## Author(s)

Marcelino de la Cruz Rot

## Examples

```
data(Astragalus)
data(Sesleria)

# Test overlapping between Astragalus individuals
# Ideally nsim should be at least 199
Astragalus.test<- test.auto.intersection(Astragalus, nsim=19)
# Observed overlapping area
Astragalus.test[1]
# p-value (negative value indicates that the observed overlapping is smaller
# than expected)
pval(Astragalus.test)

# Test overlapping between Astragalus and Sesleria individuals.
# Here, Sesleria is the accessory species (its individuals are kept fixed during the
# test) and Astragalus the focal one (its individuals are rotated)
# Ideally nsim should be at least 199
Sesleria.Astragalus.test<- test.intersection(ventana1= Sesleria,
                                              ventana2= Astragalus, nsim=19)
# Observed overlapping area
Sesleria.Astragalus.test[1]
# p-value (negative value indicates that the observed overlapping is smaller
# than expected)
pval(Sesleria.Astragalus.test)

# Reducing computing burden when making repetitive testing
```

```

# First, put all the polygonal regions in a list, i.e.
owins<- list(Astragalus, Sesleria)

# compute diameters and centroids of the individual polygons
# in each polygonal region

centroids<- list()
diams<- list()
for ( i in 1: length(owins)){
  cd<- centroiddiam(owins[[i]])
  centroids[[i]] <- cd$centroids
  diams[[i]] <- cd$diams
}

# set the number of simulations for each test
# Ideally nsim should be at least 199
online <- interactive()
Nsim <- if(online) 19 else 3

# create an array to store the results
result <- array(NA, dim=c(length(owins),length(owins),Nsim+1))

t0<-Sys.time()
for ( i in 1: length(owins)){
  for ( j in 1: length(owins)){
    cat(i,j,"\n")
    if(j!=i) result[i,j,] <- test.intersection(owins[[i]], owins[[j]], nsim=Nsim,
centroides1=centroids[[i]], diametros1=diams[[i]],
centroides2=centroids[[j]], diametros2=diams[[j]]) else
result[i,j,] <- test.auto.intersection(owins[[i]], nsim=Nsim,
centroides1=centroids[[i]], diametros1=diams[[i]])
  }
}
Sys.time()-t0

# observed values (focal species in columns)
(observed<- t(result[,,1]))
# p-values
tabla.p<- apply(result,c(1,2),pval)
(p_values <- t(tabla.p))

# Compare with parallelized versions:

# create an array to store the result.ps
result.p<- array(NA, dim=c(length(owins),length(owins),Nsim+1))

t0<-Sys.time()
for ( i in 1: length(owins)){
  for ( j in 1: length(owins)){
    cat(i,j,"\n")
    if(j!=i) result.p[i,j,] <- test.intersection.p(owins[[i]], owins[[j]], nsim=Nsim,
centroides1=centroids[[i]], diametros1=diams[[i]],

```

```
centroides2=centroids[[j]], diametros2=diams[[j]]) else
result.p[i,j,] <- test.auto.intersection.p(owins[[i]], nsim=Nsim,
centroides1=centroids[[i]], diametros1=diams[[i]])
  }
}
Sys.time()-t0

# observed values (focal species in columns)
(observed.p<- t(result.p[, ,1]))
# p-values
tabla.p.p<- apply(result.p,c(1,2),pval)
(p_values.p <- t(tabla.p.p))
```

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