

# Package ‘scp’

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**Type** Package

**Title** Spatial Conformal Prediction

**Version** 0.1.0

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**Description** Provide robust, valid, and model-free spatial prediction intervals using Spatial Conformal Prediction (SCP) algorithms

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.0.2

**URL** <https://github.com/mhuiying/scp>

**BugReports** <https://github.com/mhuiying/scp/issues>

**Suggests** knitr,  
rmarkdown,  
covr,  
testthat

**VignetteBuilder** knitr

**Imports** geoR

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krige_pred	<i>Kriging prediction function</i>
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## Description

This function provides an example for `pred_fun` in [scp](#), [plausibility](#), and [plausibility\\_contour](#), which provides a point prediction for location `s0` (and corresponding standard error), given observations `s` and `Y`.

## Usage

```
krige_pred(s0, s, Y, return_sd = FALSE)
```

## Arguments

<code>s0</code>	prediction location, a numeric vector with <code>length = 2</code> .
<code>s</code>	an $n \times 2$ matrix or <code>data.frame</code> with two coordinates of $n$ locations.
<code>Y</code>	a vector with $n$ values corresponding to $Y(s)$ .
<code>return_sd</code>	logical. if TRUE, <code>Krige_pred</code> function returns the standard error of $Y(s0)$ along with the point prediction; if FALSE, <code>Krige_pred</code> function only returns the point prediction. Defaults to FALSE.

## Value

The output is a value of point prediction for  $Y(s0)$  if `return_sd` is FALSE or a list with the following elements if `return_sd` is TRUE.

<code>yhat</code>	point prediction for $Y(s0)$
<code>sd</code>	standard error for $Y(s0)$

## Examples

```
N = 21; n = N^2
S = seq(0,1,length=N)
s = expand.grid(S,S)
d = as.matrix(dist(s))

theta      = c(0,3,0.1,0.7)
names(theta) = c("Nugget","PartialSill","Range","Smoothness")
C = mat_cov(d,theta)
X = t(chol(C))%*%rnorm(n)
Y = X^3 + rnorm(n)

s0 = c(0.5, 0.5)
krige_pred(s0,s,Y)
krige_pred(s0,s,Y,interval=TRUE)
```

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plausibility	<i>calculate plausibility for <math>Y_0</math></i>
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## Description

This function provides the plausibility of  $Y(s_0)$  being  $Y_0$ , given observations  $s$  and  $Y$ , using spatial conformal prediction algorithms.

## Usage

```
plausibility(
  Y0,
  s0,
  s,
  Y,
  global = TRUE,
  eta = Inf,
  m = NULL,
  pred_fun = krige_pred,
  dfun = c("residual2", "abs_residual", "std_residual2", "std_abs_residual")
)
```

## Arguments

$Y_0$	a scalar or a vector
$s_0$	prediction location, a numeric vector with <code>length = 2</code> .
$s$	an $n \times 2$ matrix or data.frame with two coordinates of $n$ locations.
$Y$	a vector with $n$ values corresponding to $Y(s)$ .
<code>global</code>	logical; if <code>TRUE</code> , <code>scp</code> function returns the result of global spatial conformal prediction (GSCP); if <code>FALSE</code> , <code>scp</code> function returns the result of local spatial conformal prediction (LSCP) and users need to specify <code>eta &lt; Inf</code> or <code>m ≤ n</code> . Defaults to <code>TRUE</code> .
<code>eta</code>	kernel bandwidth for weight schema, a positive scalar with smaller value meaning more localized procedure. Defaults to <code>Inf</code> , which puts equal weight on surrounding <code>m</code> points.
<code>m</code>	an positive integer representing the number of nearest locations to use for prediction. Default to <code>NULL</code> . If <code>global = TRUE</code> , <code>m = n</code> ; if <code>global = FALSE</code> and <code>m</code> is not specified, <code>m</code> would be determined by <code>eta</code> .
<code>pred_fun</code>	spatial prediction function with inputs being <code>s0,s,Y</code> and outputs being predicted $Y(s_0)$ (and its standard error). Defaults to <a href="#">krige_pred</a> .
<code>dfun</code>	non-conformity measure with four options. In which, <code>"residual2"</code> (default) represents squared residual, <code>"std_residual2"</code> represents standardized squared residual, <code>"abs_residual"</code> represents absolute residual, and <code>"std_abs_residual"</code> represents standardized absolute residual.

**Value**

The output is a scalar or a vector with plausibility values for  $Y_0$ . The numbers are between 0 and 1.

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**References**

to be entered

**See Also**

[plausibility\\_contour](#)

**Examples**

```
N = 41; n = N^2
S = seq(0,1,length=N)
s = expand.grid(S,S)
d = as.matrix(dist(s))

theta      = c(0,3,0.1,0.7)
names(theta) = c("Nugget", "PartialSill", "Range", "Smoothness")

C = mat_cov(d,theta)
X = t(chol(C))%*%rnorm(n)
Y = X^3 + rnorm(n)

s0 = c(0.5, 0.5)
idx = which(s[,1]==s0[1] & s[,2]==s0[2])

plausibility(Y0=seq(0,1,0.1),s0=s0,s=s[-idx,],Y=Y[-idx])
```

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plausibility\_contour    *generate plausibility contour*

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**Description**

This function provides the plausibility contour for  $Y(s_0)$ , given observations  $s$  and  $Y$ , using spatial conformal prediction algorithms.

## Usage

```
plausibility_contour(
  s0,
  s,
  Y,
  global = TRUE,
  eta = Inf,
  m = NULL,
  pred_fun = krige_pred,
  dfun = c("residual2", "abs_residual", "std_residual2", "std_abs_residual"),
  precision = NULL
)
```

## Arguments

<code>s0</code>	prediction location, a numeric vector with <code>length = 2</code> .
<code>s</code>	an $n \times 2$ matrix or data.frame with two coordinates of $n$ locations.
<code>Y</code>	a vector with $n$ values corresponding to $Y(s)$ .
<code>global</code>	logical; if TRUE, <code>scp</code> function returns the result of global spatial conformal prediction (GSCP); if FALSE, <code>scp</code> function returns the result of local spatial conformal prediction (LSCP) and users need to specify <code>eta &lt; Inf</code> or $m \leq n$ . Defaults to TRUE.
<code>eta</code>	kernel bandwidth for weight schema, a positive scalar with smaller value meaning more localized procedure. Defaults to Inf, which puts equal weight on surrounding $m$ points.
<code>m</code>	an positive integer representing the number of nearest locations to use for prediction. Default to NULL. If <code>global = TRUE</code> , $m = n$ ; if <code>global = FALSE</code> and $m$ is not specified, $m$ would be determined by <code>eta</code> .
<code>pred_fun</code>	spatial prediction function with inputs being <code>s0, s, Y</code> and outputs being predicted $Y(s0)$ (and its standard error). Defaults to <a href="#">krige_pred</a> .
<code>dfun</code>	non-conformity measure with four options. In which, "residual2" (default) represents squared residual, "std_residual2" represents standardized squared residual, "abs_residual" represents absolute residual, and "std_abs_residual" represents standardized absolute residual.
<code>precision</code>	a positive scalar represents how dense $Y(s)$ candidates ( <code>Y_cand</code> ) are. Defaults to NULL.

## Value

The output is a data.frame of `Y_cand` and corresponding plausibility values

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## References

to be entered

**See Also**[plausibility](#)


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 scp
 

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*Spatial conformal prediction at a single input location*


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**Description**

This function provides the spatial conformal prediction interval for location `s0`, given observations `s` and `Y`.

**Usage**

```
scp(
  s0,
  s,
  Y,
  global = TRUE,
  eta = Inf,
  m = NULL,
  pred_fun = krige_pred,
  dfun = c("residual2", "abs_residual", "std_residual2", "std_abs_residual"),
  precision = NULL,
  alpha = 0.05
)
```

**Arguments**

<code>s0</code>	prediction location, a numeric vector with <code>length = 2</code> .
<code>s</code>	an $n \times 2$ matrix or <code>data.frame</code> with two coordinates of $n$ locations.
<code>Y</code>	a vector with $n$ values corresponding to $Y(s)$ .
<code>global</code>	logical; if <code>TRUE</code> , <code>scp</code> function returns the result of global spatial conformal prediction (GSCP); if <code>FALSE</code> , <code>scp</code> function returns the result of local spatial conformal prediction (LSCP) and users need to specify <code>eta</code> . Defaults to <code>TRUE</code> .
<code>eta</code>	kernel bandwidth for weight schema, a positive scalar with smaller value meaning more localized procedure. Defaults to <code>Inf</code> , which puts equal weight on surrounding $m$ points.
<code>m</code>	an positive integer representing the number of nearest locations to use for prediction. Default depends on <code>eta</code> .
<code>pred_fun</code>	spatial prediction function with inputs being <code>s0, s, Y</code> and outputs being predicted $Y(s0)$ (and its standard error). Defaults to <a href="#">krige_pred</a> representing Kriging prediction.

<code>dfun</code>	non-conformity measure with four options. In which, <code>"residual2"</code> (default) represents squared residual, <code>"std_residual2"</code> represents standardized squared residual, <code>"abs_residual"</code> represents absolute residual, and <code>"std_abs_residual"</code> represents standardized absolute residual.
<code>precision</code>	a positive scalar represents how dense the candidates for $Y(s)$ are. Defaults to NULL.
<code>alpha</code>	significance level. Defaults to 0.05.

### Value

The output is a vector of lower and upper bounds of the conformal prediction interval.

### Author(s)

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### References

to be entered

### See Also

[plausibility](#)

### Examples

```

N = 41; n = N^2
S = seq(0,1,length=N)
s = expand.grid(S,S)
d = as.matrix(dist(s))

theta      = c(0,3,0.1,0.7)
names(theta) = c("Nugget","PartialSill","Range","Smoothness")
C = mat_cov(d,theta)
X = t(chol(C))%*%rnorm(n)
Y = X^3 + rnorm(n)

s0 = c(0.5, 0.5)
idx = which(s[,1]==s0[1] & s[,2]==s0[2])
pred_fun = function(s0,s,Y,alpha=0.05) return(mean(Y))
PI = scp(s0,s[-idx,],Y[-idx],pred_fun=pred_fun, dfun="abs_residual")
cat(paste("True value: ", Y[idx], "\n"))
cat(paste("Prediction Interval: [ ", PI[1], ", ", PI[2], " ]"))

```

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