# Package 'sdfEXTREME'

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Title Spatial Deformation for Non-Stationary Extremal Dependence

Type Package

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station	Contains functions for creating spatial deformations to account for non- arity in spatial data. Both correlation based methods and methods adapted for extremal de- ice are included. Accompanies the paper of the same title.
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NeedsComp	
R topics	documented:
Aus	s_Heat
Aus	s_Heat_Output
brns	sims
	3
	S.emp
	dSplineParFNCOR
	dSplineParFNEXTR
_	nsv
_	h_Precip_Output
-	eject
	tepairs
	IT
nllI	MSPexp
nllI	MSPexpSmith
nllN	MSPexp
retu	rnDcoord
Sco	re_Est

2 Aus\_Heat\_Output

Index		29
	Vterm	28
	stat.boot	27
	Snow_Precip_Output	26
	Snow_Precip	26
	sdf.heur.EXTR	24
	sdf.heur.Cor	22

Aus\_Heat

Australia Summer Temperatures

# **Description**

Data consist of daily summer (DJF) maximum near-surface air temperatures taken from the HadGHCND global gridded dataset and interpolated to 72 grid point locations covering Australia, for the period 1957-2014.

#### Usage

```
data(Aus_Heat)
```

#### **Format**

A list with 2 elements:

**Temp.** A matrix with 5234 rows and 72 columns of temperature data.

coords A 72 by 2 matrix of lon-lat coordinates.

# References

```
Caesar et al. (2006) J. Geophys. Res. 111, D05101, (doi)
```

# **Examples**

```
data(Aus_Heat)
```

Aus\_Heat\_Output

Australia Summer Temperatures outputs

# Description

Outputs from deformations and model fits for data(Aus\_Heat).

# Usage

```
data(Aus_Heat_Output)
```

brnsims 3

#### **Format**

A list with 5 elements:

**likG.IMSP** Optim output from fitting Inverted Smith model to G-plane sampling locations. See nllIMSPexpSmith.

**likG.MSP** Optim output from fitting Brown-Resnick model to G-plane sampling locations. See nllMSPexpSmith.

**likD.IMSP** Optim output from fitting Inverted Smith model to D-plane sampling locations. See nllIMSPexp.

**likD.MSP** Optim output from fitting Brown-Resnick model to D-plane sampling locations. See nllMSPexp.

sdf D-plane transformation spline parameters. See sdf.heur.EXTR.

#### References

```
Caesar et al. (2006) J. Geophys. Res. 111, D05101, (doi)
```

# **Examples**

```
data(Aus_Heat_Output)
```

brnsims

Simulate non-stationary Brown-Resnick process

#### **Description**

Simulate a non-stationary (or stationary) Brown-Resnick process using the 'stopping rule' methodology of Dieker and Mikosch (2015). Non-stationary Brown-Resnick processes are simulated using the non-stationary semivariogram. detailed in the help file for gamsv.

# Usage

```
brnsims(reps, locs, kappa, lambda, tau, centre = NULL)
```

#### **Arguments**

reps Number of realisations, n. locs A p by 2 matrix of coordinates. kappa Value of  $\kappa$ . See help(gamsv). lambda Value of  $\lambda$ . See help(gamsv).

tau A p by p matrix of correlation values.

centre Vector of length 2 giving coordinates of centre of non-stationarity for semivari-

ogram. If NULL, stationary semivariogram used. See help(gamsv).

# Value

n by p matrix

4 chi3

#### References

Dieker and Mikosch (2015) Extremes, 18(2):301-314, (doi)

#### **Examples**

```
##Creating correlation values to simulate non-stationary Brown-Resnick process.
lambda<-2
centre < -c(0,0)
kappa<-0.8
n.grid<-8
sim.coords<-as.matrix(expand.grid(seq(-1,1,length=n.grid)),seq(-1,1,length=n.grid)))</pre>
p<-dim(sim.coords)[1]</pre>
tau<-matrix(NA,nrow=p,ncol=p)</pre>
for(i in 1:p){
for(j in 1:p){
tau[i,j] <- gamsv(s1=sim.coords[i,],s2=c(0,0),lam=lambda,kap=kappa,centre=NULL) + lambda,kap=kappa,centre=NULL) + lambda,kappa,centre=NULL) + lambda
gamsv(s1=sim.coords[j,],s2=c(0,0),lam=lambda,kap=kappa,centre=NULL)-
gamsv(s1=sim.coords[i,],s2=sim.coords[j,],lam=lambda,kap=kappa,centre=centre)
}
}
\#\#Simulates 10 realisations of non-stationary BR process.
\verb|Sim<-braims(reps=10,locs=sim.coords,kappa=kappa,lambda=lambda,centre=centre,tau=tau)|
```

chi3

Theoretical triple-wise  $\chi$  ( $\chi$ \_q) for a (Inverted) Brown-Resnick process

#### **Description**

Calculates the theoretical triple-wise  $\chi(s_i, s_j, s_k)$  or  $\chi_q(s_i, s_j, s_k)$  for a given 3 by 3 matrix of semivariogram. values.

#### Usage

```
chi3(v_H)
chi3q(v_H, q)
```

# Arguments

v\_H 3 by 3 matrix of semivariogram. values. q Exceedance threshold for  $\chi_q(s_i,s_j,s_k)$ .

chi3.emp 5

#### Value

Theoretical  $\chi(s_i, s_j, s_k)$  or  $\chi_q(s_i, s_j, s_k)$  measure for the corresponding matrix of semivariogram values.

# **Examples**

```
data(Aus_Heat)
data(Aus_Heat_Output)
Z<-Aus_Heat$Temp.
Gcoords<-Aus_Heat$coords
sdf<-Aus_Heat_Output$sdf
likD.MSP<-Aus_Heat_Output$likD.MSP
likD.IMSP<-Aus_Heat_Output$likD.IMSP

Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)

ind.triple<-c(1,2,3) ##Denotes indices of triple for which triple-wise chi calculated.
#MSP
v_H<-(rdist.earth(Dcoords[ind.triple,],miles=F)/likD.MSP$par[2])^likD.MSP$par[1]
print(chi3(v_H))

#IMSP - likD.IMSP only contains range parameter as smoothing parameter is 2.
v_H<-(rdist.earth(Dcoords[ind.triple,],miles=F)/likD.IMSP$par[1])^2
print(chi3q(v_H,q=0.95))</pre>
```

chi3.emp

Empirical estimate of triple-wise chi

# Description

Calculates empirical estimate of triple-wise chi for (U, V, W).

# Usage

```
chi3.emp(U, V, W, q)
```

# **Arguments**

U	Vector of length $n$ of standard uniform variables.
٧	Vector of length $n$ of standard uniform variables.
W	Vector of length $n$ of standard uniform variables.
q	Value of exceedance probability used in estimation.

# Value

Estimate of triple-wise chi for (U, V, W).

# **Examples**

```
data(Aus_Heat)
Z<-Aus_Heat$Temp.
unif<-function(x) rank(x)/(length(x)+1)
Z_U<-Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
q<-0.95
ind.triple<-c(1,2,3) ##Denotes indices of triple for which triple-wise chi calculated.
chi3.emp(Z_U[,ind.triple[1]],Z_U[,ind.triple[2]],Z_U[,ind.triple[3]],q)</pre>
```

FindSplineParFNCOR

Find spline parameters (Correlation)

# **Description**

Find spline parameters for deformation based on correlation methods. Either use the Frobenuis norm via type=="F-norm" or the original Smith (1996) method via type=="Smith". Output to be minimised to estimate spline parameters.

# Usage

```
FindSplineParFNCOR(
  par,
  Gcoords,
  m.ind,
  n = 0,
  emp.cor,
  type = c("F-norm", "Smith"),
  sphere.dis = FALSE
)
```

# **Arguments**

par	Parameter values $\phi = (b_1, b_2, \rho, \kappa, \delta_4^{(1)}\delta_m^{(1)}, \delta_4^{(2)}\delta_m^{(2)})$ . If $m < 4$ , $\delta$ parameters are not needed.
Gcoords	A d by 2 matrix of G-plane coordinates.
m.ind	A vector of length m <d anchor="" gcoords.<="" giving="" in="" indices="" of="" points="" td="" the=""></d>
n	Number of data points used to estimate emp. cor. Only necessary for type=="Smith".
emp.cor	A d by d matrix of pairwise empirical correlation values.
type	"F-norm" for Frobenius norm method using theoretical $\rho(h_{ij}^*)$ from a stationary Matérn correlation model. "Smith" for original Smith (1996) method; also uses stationary Matérn correlation model.
sphere.dis	Is Spherical distance or Euclidean distance used?

#### Value

Frobenius norm of difference between theoretical  $\rho(h_{ij}^*)$  matrix and emp.cor, or negative log-likelihood for a stationary Gaussian process fit using D-plane coordinates.

```
data("Aus_Heat")
Z<-Aus_Heat$Temp.
Gcoords<-Aus_Heat$coords
Z_U < -Z
unif<-function(x){rank(x)/(length(x)+1)}</pre>
#Transform to uniform margins
for(i in 1:dim(Z_U)[2]){
  Z_U[,i] < -unif(Z[,i])
#Transform to Gaussian margins
Z_N<-qnorm(Z_U)</pre>
#Calculate pairwise empirical correlation
emp.cor<-matrix(rep(0, dim(Z_N)[2]^2), nrow=dim(Z_N)[2], ncol=dim(Z_N)[2])
for(i in 1:dim(Z_N)[2]){
  for(j in 1:i){
    emp.cor[i,j] < -cor(Z_N[,i],Z_N[,j])
  }
}
emp.cor<-emp.cor+t(emp.cor)</pre>
diag(emp.cor)<-diag(emp.cor)/2</pre>
# Set number of anchor points
m < -10
# Sample anchor points
m.ind<-sample(1:dim(Gcoords)[1],m,replace=FALSE)</pre>
#Transform to D-plane using Frobenius norm method
sdf < -optim(fn=FindSplineParFNCOR, par=c(0.05, 0.05, 0, 1, rep(0, 2*m-6)),
           type="F-norm", sphere.dis=TRUE, Gcoords=Gcoords, m.ind=m.ind,
           control=list(maxit=2000), emp.cor=emp.cor,method = "Nelder-Mead")
sdf<-try(optim(fn=FindSplineParFNCOR,par=sdf$par,sphere.dis=TRUE,type="F-norm",
         Gcoords=Gcoords,m.ind=m.ind, control=list(maxit=2000), emp.cor=emp.cor,
    method = "BFGS"))
sdf$m.ind<-m.ind
#Plot Dcoords
Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)
plot(Dcoords)
```

FindSplineParFNEXTR Find spline parameters (Extremal)

# **Description**

Find spline parameters for deformation using extremal dependence methods. Provides Frobenuis norm of difference between pairwise empircal and theoretical  $\chi$  measures. Output to be minimised to estimate spline parameters.

# Usage

```
FindSplineParFNEXTR(
  par,
  Gcoords,
  m.ind,
  emp.dep,
  type = c("CHI_BR", "CHI_q_IBR"),
  q = 0,
  sphere.dis = FALSE
)
```

# Arguments

par	Parameter values $\phi=(b_1,b_2,\rho,\kappa,\delta_4^{(1)}\delta_m^{(1)},\delta_4^{(2)}\delta_m^{(2)}).$ If m < 4, $\delta$ parameters are not needed.
Gcoords	A d by 2 matrix of G-plane coordinates.
m.ind	A vector of length $m < d$ giving the indices of the anchor points in Gcoords.
emp.dep	A d by d matrix of pairwise empirical chi values.
type	"CHI_BR" for theoretical $\chi(h_{ij}^*)$ from a Brown-Resnick model. "CHI_q_IBR" for theoretical $\chi_q(h_{ij}^*)$ from an inverted Brown-Resnick model.
q	Threhold for $\chi_q(h_{ij}^*)$ . Only needed if type=="CHI_q_IBR".
sphere.dis	Is Spherical distance or Euclidean distance used?

# Value

Frobenius norm of difference between theoretical  $\chi(h_{ij}^*)$  or  $\chi_q(h_{ij}^*)$  matrix and emp.chi.

```
data("Aus_Heat")
Z<-Aus_Heat$Temp.
Gcoords<-Aus_Heat$coords

Z_U<-Z
unif<-function(x){rank(x)/(length(x)+1)}
#Transform to uniform margins
for(i in 1:dim(Z_U)[2]){</pre>
```

gamsv 9

```
Z_U[,i] < -unif(Z[,i])
\label{eq:chi.emp} \begin{split} \text{chi.emp} &-\text{function}(\mathsf{U},\mathsf{V},\mathsf{z}) \ \text{sum}((\mathsf{U} \!\!> \!\! \mathsf{z}) \& (\mathsf{V} \!\!> \!\! \mathsf{z})) / \text{sum}(\mathsf{U} \!\!> \!\! \mathsf{z}) \end{split}
q < -0.98
#Calculate pairwise empirical chi
emp.chi<-matrix(rep(0,dim(Z_U)[2]^2),nrow=dim(Z_U)[2],ncol=dim(Z_U)[2])
for(i in 1:dim(Z_U)[2]){
      for(j in 1:i){
             emp.chi[i,j] < -chi.emp(Z_U[,i],Z_U[,j],q)
       }
}
emp.chi<-emp.chi+t(emp.chi)</pre>
diag(emp.chi)<-diag(emp.chi)/2</pre>
# Set number of anchor points
m<-10
# Sample anchor points
m.ind<-sample(1:dim(Gcoords)[1],m,replace=FALSE)</pre>
#Transform to D-plane using Brown-Resnick theoretical chi function
sdf < -optim(fn = FindSplineParFNEXTR, par = c(0.05, 0.05, 0, 1, rep(0, 2 *m - 6)), type = "CHI\_BR", sphere.dis = TRUE, for the substitution of 
                                    Gcoords=Gcoords,m.ind=m.ind,control=list(maxit=2000), emp.dep=emp.chi,
                    method = "Nelder-Mead")
sdf<-optim(fn=FindSplineParFNEXTR,par=sdf$par,sphere.dis=TRUE,type="CHI_BR",</pre>
                                    Gcoords=Gcoords, m.ind=m.ind,
                                    control=list(maxit=2000), emp.dep=emp.chi,method = "BFGS")
 sdf$m.ind<-m.ind
#Plot Dcoords
Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)
plot(Dcoords)
```

gamsv

Non-stationary 'centered' semivariogram.

# **Description**

Calculates the Fouedijo et al. (2015) theoretical semivariogram. This is a function of some central location o = centre and distance between locations s1 and s2. The semivariogram. has the form

$$\gamma^*(s_2,s_1)=\gamma(||\psi(s_2)-\psi(s_1)||),$$
 where 
$$\psi(s)=o+(s-o)||s-o||$$
 and 
$$\gamma(x)=(x/\lambda)^\kappa$$
 for  $\lambda>0$  and  $\kappa\in(0,2].$ 

10 gamsv

#### Usage

```
gamsv(s1, s2, centre = NULL, lam, kap)
```

# **Arguments**

Vector of length 2 giving coordinates of first location. Vector of length 2 giving coordinates of second location. Vector of length 2 giving coordinates of centre of non-stationarity. If NULL, returns stationary semivariogram. Value of  $\lambda$ .

kap Value of  $\kappa$ .

#### Value

Non-stationary semivariogram. between locations s1 and s2

#### References

Fouedijo et al. (2015) Spatial Statistics, 13:45-61, (doi)

```
##Creating correlation values to simulate non-stationary Brown-Resnick process. See help(brnsims).
lambda<-2
centre < -c(0,0)
kappa<-0.8
n.grid<-8
sim.coords<-as.matrix(expand.grid(seq(-1,1,length=n.grid),seq(-1,1,length=n.grid)))</pre>
p<-dim(sim.coords)[1]</pre>
tau<-matrix(NA,nrow=p,ncol=p)</pre>
for(i in 1:p){
for(j in 1:p){
tau[i,j] < -gamsv(s1=sim.coords[i,],s2=c(0,0),lam=lambda,kap=kappa,centre=NULL) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0,0) + (0
{\tt gamsv(s1=sim.coords[j,],s2=c(0,0),lam=lambda,kap=kappa,centre=NULL)-}
gamsv(s1=sim.coords[i,],s2=sim.coords[j,],lam=lambda,kap=kappa,centre=centre)
}
}
##Simulates 10 realisations of non-stationary BR process.
\verb|Sim<-braims(reps=10,locs=sim.coords,kappa=kappa,lambda=lambda,centre=centre,tau=tau)|
```

High\_Precip 11

High\_Precip

Highlands Precipitation Data

# Description

Data consist of winter (DJF) 12-hour average precipitation rate (mm/day) taken from the UKCP18 climate projection. Data is produced on  $2.2 \times 2.2 km^2$  grid-boxes, between the years 1980 and 2000.

# Usage

```
data(High_Precip)
```

#### **Format**

A list with 2 elements:

Pr. A matrix with 3600 rows and 100 columns of precipitation data.

coords A 100 by 2 matrix of lon-lat coordinates, corresponding to the centroid of each grid-box.

#### References

```
Lowe et al. (2018), Met Office, Exter, UK, (pdf)
```

#### **Examples**

```
data(High_Precip)
```

High\_Precip\_Output

Highlands Precipitation outputs

# Description

Outputs from deformations and model fitting for data(High\_Precip).

#### Usage

```
data(High_Precip_Output)
```

#### **Format**

A list with 5 elements:

- **likG.IMSP** Optim output from fitting Inverted Brown-Resnick model to G-plane sampling locations. See nllIMSPexp.
- **likG.MSP** Optim output from fitting Brown-Resnick model to G-plane sampling locations. See nllMSPexp.
- **likD.IMSP** Optim output from fitting Inverted Brown-Resnick model to D-plane sampling locations. See nllIMSPexp.
- **likD.MSP** Optim output from fitting Brown-Resnick model to D-plane sampling locations. See nllMSPexp.
- sdf D-plane transformation spline parameters. See sdf.heur.EXTR.

m.reject

#### References

```
Lowe et al. (2018) Met Office, Exter, UK, (pdf)
```

# **Examples**

```
data(High_Precip_Output)
```

m.reject

A heuristic for deciding if initial anchor points have enough spread.

# **Description**

This heuristic ensures that chosen anchor points are not along a straight line and, if length(m.ind)<5, that the maximum distance between the initial anchor points in the G-plane is sufficiently large i.e. over 60% of the maximum pairwise distance of all sampling locations in the G-plane.

# Usage

```
m.reject(m.ind, Gcoords, sphere.dis = FALSE)
```

# **Arguments**

m. ind A vector of indices denoting the anchor points in Gcoords.

Gcoords A d by 2 matrix of G-plane sampling locations.

sphere.dis Is Spherical distance or Euclidean distance used?

# Value

1 if anchor points are rejected, 0 otherwise.

```
data("Aus_Heat")
Gcoords<-Aus_Heat$coords
# Set number of anchor points
m<-10
# Sample anchor points
m.ind<-sample(1:dim(Gcoords)[1],m,replace=FALSE)
reject<-m.reject(m.ind,Gcoords,sphere.dis=TRUE)
print(reject)</pre>
```

makepairs 13

makepairs	Make pairs
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# **Description**

Produce a list of all configurations of pairwise exceedances above some threshold u. Used in censored likelihood estimation.

#### Usage

```
makepairs(Z, u)
```

# **Arguments**

Z A N by d matrix.

u Censoring threshold.

#### Value

List of four data frames. Each data frame has 4 columns and the rows sum to N\*d\*(d-1)/2. For each data frame the first two columns are the values and the last two are the indices for the location of these values in the columns of Z.

Component 1 Pairs of observations where both components exceed u.

Component 2 Pairs of observations where the first component exceeds u and the second equals u.

Component 3 Pairs of observations where the second component exceeds u and the first equals u.

**Component 4** Pairs of observations where both components equal u.

```
#For a N by d matrix of data "Z" and d by 2 matrix of coordinates "Gcoords".
# We use a very small subset of data(Aus_Heat) as an example.
##THIS WILL TAKE A LONG TIME TO RUN WITH THE FULL DATASET##

data(Aus_Heat)
Z<-Aus_Heat$Temp.[,1:3]
Gcoords<-Aus_Heat$coords[1:3,]

unif<-function(x) rank(x)/(length(x)+1)
Z_U<-Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
Z_Exp<-qexp(Z_U) #Transform to exponential margins

q<-0.98
u<-quantile(Z_Exp,prob=q) # Censoring threshold
# Create a list of length 4 of pairwise exceedances and index in coordinate matrix
Zpair<-makepairs(Z_Exp,u=u)</pre>
```

14 nllHT

nllHT Negative log-likelihood for bivariate Heffernan and To model	Tawn (2004)
-----------------------------------------------------------------------	-------------

# Description

Calculates the negative log-likelihood for Y|X=x. This is a profile log-likelihood with free parameters  $\alpha$  and  $\beta$ . We use the same normalising functions as in the original paper.

#### Usage

```
nllHT(X, Y, par)
```

# Arguments

```
X Vector of Laplace variables. This is the conditioning variable i.e. X=x. Y Vector of Laplace variables, Y. Vector (\alpha,\beta).
```

#### Value

Negative log-likelihood for Heffernan and Tawn model fit to Y|X=x.

#### References

Heffernan and Tawn (2004), Journal of the Royal Statistical Society: Series B, 66:497-546, (doi)

```
#For a N by d matrix of data "Z" and d by 2 matrix of coordinates "Gcoords".
# We use data(Aus_Heat) as an example.
library(rmutil)
library(fields)
data(Aus_Heat)
Z<-Aus_Heat$Temp.</pre>
Gcoords<-Aus_Heat$coords
unif<-function(x) rank(x)/(length(x)+1)
Z_U < -Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
Z_LP<-qlaplace(Z_U) # Transform to Laplace margins</pre>
p<-dim(Gcoords)[1]</pre>
ConExp<-matrix(0,nrow=p,ncol=p)</pre>
u<-quantile(Z_LP,0.98) \#Quantile for estimating conditional expectation
#Calculate conditional expectation for each pair
for(i in 1:p){
 for(j in 1:i){
   Exceedances<-cbind(Z_LP[,i],Z_LP[,j])[which(Z_LP[,i]>=u),]
```

nlIIMSPexp 15

```
opt<-optim(nllHT,X=Exceedances[,1],Y=Exceedances[,2],par=c(0.3,0.8))
    #print(opt)
    alpha<-opt$par[1]
    beta<-opt$par[2]
    mu<-mean((Exceedances[,2]-alpha*Exceedances[,1])/Exceedances[,1]^beta)
    ConExp[i,j]<-alpha*u+u^beta*mu
}

ConExp<-ConExp+t(ConExp)  ##Symmetry assumed
diag(ConExp)<-diag(ConExp)/2

plot(rdist.earth(Gcoords,miles=F),ConExp, ylab="Conditional Expectation",
    xlab="Distance (Km)", main="")</pre>
```

nllIMSPexp Negative composite censored log-likelihood for the stationary inverted Brown-Resnick process

# **Description**

Calculate the negative composite censored log-likelihood for the stationary inverted Brown-Resnick process on standard exponential margins.

#### Usage

```
nllIMSPexp(par, ZBE, ZXE, ZYE, ZNE, coord, n.a, sphere.dis = F)
```

#### **Arguments**

par	Stationary semivariogram parameters $(\kappa, \lambda)$ .
ZBE	A list of length d with each element a matrix with 2 columns.
ZXE	A list of length d with each element a matrix with 2 columns.
ZYE	A list of length d with each element a matrix with 2 columns.
ZNE	A list of length d with each element a matrix with 2 columns.
coord	A d by 2 matrix of coordinates.
n.a	A vector with length determined by the number of unique pairs in ZNE.
sphere.dis	Is Spherical distance or Euclidean distance used?

# Value

Negative composite censored log-likelihood for inverted Brown-Resnick process.

```
# For a N by d matrix of data "Z" and d by 2 matrix of coordinates "Gcoords".
# We use a very small subset of data(Aus_Heat) as an example.
##THIS WILL TAKE A LONG TIME TO RUN WITH THE FULL DATASET##
library(fields)
data(Aus_Heat)
```

16 nllIMSPexpSmith

```
Z<-Aus_Heat$Temp.[,1:3]</pre>
Gcoords<-Aus_Heat$coords[1:3,]</pre>
unif<-function(x) rank(x)/(length(x)+1)
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
Z_Exp < -qexp(Z_U) #Transform to exponential margins
a < -0.98
u<-quantile(Z_Exp,prob=q) # Censoring threshold
# Create a list of length 4 of pairwise exceedances and index in coordinate matrix
Zpair<-makepairs(Z_Exp,u=u)</pre>
#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))</pre>
dist2<-apply(Zpair[[4]][,3:4],1,function(x){dist[x[1],x[2]]})</pre>
unique.d<-as.matrix(unique(dist2))</pre>
n.a<-rep(0, dim(unique.d)[1])</pre>
for(i in 1:length(unique.d)){
 n.a[i]<-sum(dist2==unique.d[i,1])</pre>
}
#WARNING- pairwise likelihoods take some time to run
\# Inverted Brown-Resnick process fit to G-plane coordinate system
likG.IMSP<-optim(fn=nllIMSPexp,par=c(1,500),ZBE=as.matrix(Zpair[[1]]),</pre>
           ZXE=as.matrix(Zpair[[2]]),
           ZYE=as.matrix(Zpair[[3]]), ZNE=as.matrix(Zpair[[4]]),
           n.a=n.a,sphere.dis=TRUE,coord=Gcoords,
           control=list(maxit=2000),
           method = "Nelder-Mead",hessian=TRUE)
```

nllIMSPexpSmith

Smith process likelihoods

# **Description**

Calculate the negative composite censored log-likelihood for the Smith and Inverted Smith models on standard exponential margins.

#### Usage

```
nllIMSPexpSmith(par, ZBE, ZXE, ZYE, ZNE, coord, n.a, sphere.dis = F)
nllMSPexpSmith(par, ZBE, ZXE, ZYE, ZNE, coord, n.a, sphere.dis = F)
```

#### **Arguments**

par	Stationary semivariogram parameter $\lambda$ .
ZBE	A list of length d with each element a matrix with 2 columns.
ZXE	A list of length d with each element a matrix with 2 columns.
ZYE	A list of length d with each element a matrix with 2 columns.
ZNE	A list of length d with each element a matrix with 2 columns.

nllMSPexp 17

coord A d by 2 matrix of coordinates.

n.a A vector with length determined by the number of unique pairs in ZNE.

sphere.dis Is Spherical distance or Euclidean distance used?

#### Value

Negative composite censored log-likelihood for Smith (or inverted Smith) model.

# Examples

```
# For a N by d matrix of data "Z" and d by 2 matrix of coordinates "Gcoords". We use a very
# small subset of data(Aus_Heat) as an example.
##THIS WILL TAKE A LONG TIME TO RUN WITH THE FULL DATASET##
library(fields)
data(Aus_Heat)
Z<-Aus_Heat$Temp.[,1:3]</pre>
Gcoords<-Aus_Heat$coords[1:3,]</pre>
unif<-function(x) rank(x)/(length(x)+1)</pre>
Z_U < -Z
for
(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) \# Transform to uniform margins
Z_Exp<-qexp(Z_U) #Transform to exponential margins
q<-0.98
u<-quantile(Z_Exp,prob=q) # Censoring threshold
# Create a list of length 4 of pairwise exceedances and index in coordinate matrix
Zpair<-makepairs(Z_Exp,u=u)</pre>
#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))</pre>
dist2<-apply(Zpair[[4]][,3:4],1,function(x){dist[x[1],x[2]]})</pre>
unique.d<-as.matrix(unique(dist2))</pre>
n.a<-rep(0, dim(unique.d)[1])</pre>
for(i in 1:length(unique.d)){
  n.a[i]<-sum(dist2==unique.d[i,1])</pre>
\#WARNING- pairwise likelihoods take some time to run
# Inverted Smith process fit to G-plane coordinate system
likG.INV.SMITH<-optim(fn=nllIMSPexpSmith,par=c(1000),lower=0,upper=2000,ZBE=as.matrix(Zpair[[1]]),
           ZXE=as.matrix(Zpair[[2]]),
           ZYE=as.matrix(Zpair[[3]]), ZNE=as.matrix(Zpair[[4]]),
           n.a=n.a, sphere.dis=TRUE, coord=Gcoords,
           control=list(maxit=2000),
           method = "Brent",hessian=TRUE)
```

nllMSPexp

Negative composite censored log-likelihood for the stationary Brown-Resnick process 18 nllMSPexp

#### **Description**

Calculate the negative composite censored log-likelihood for the stationary Brown-Resnick process on standard exponential margins.

# Usage

```
nllMSPexp(par, ZBE, ZXE, ZYE, ZNE, coord, n.a, sphere.dis = F)
```

# **Arguments**

Stationary semivariogram parameters $(\kappa, \lambda)$ .
A list of length d with each element a matrix with 2 columns.
A list of length d with each element a matrix with 2 columns.
A list of length d with each element a matrix with 2 columns.
A list of length d with each element a matrix with 2 columns.
A d by 2 matrix of coordinates.
A vector with length determined by the number of unique pairs in ZNE.
Is Spherical distance or Euclidean distance used?

# Value

Negative composite censored log-likelihood for Brown-Resnick process.

```
\mbox{\# For a N by d matrix of data "Z" and d by 2 matrix of coordinates "Gcoords".}
# We use a very small subset of data(Aus_Heat) as an example.
##THIS WILL TAKE A LONG TIME TO RUN WITH THE FULL DATASET##
library(fields)
data(Aus_Heat)
Z<-Aus_Heat$Temp.[,1:3]</pre>
Gcoords<-Aus_Heat$coords[1:3,]</pre>
unif<-function(x) rank(x)/(length(x)+1)
Z_U < -Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
Z_Exp < -qexp(Z_U) #Transform to exponential margins
q<-0.98
u<-quantile(Z_Exp,prob=q) # Censoring threshold
# Create a list of length 4 of pairwise exceedances and index in coordinate matrix
Zpair<-makepairs(Z_Exp,u=u)</pre>
#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))</pre>
dist2 <-apply(Zpair[[4]][,3:4],1,function(x)\{dist[x[1],x[2]]\})
unique.d<-as.matrix(unique(dist2))</pre>
n.a<-rep(0, dim(unique.d)[1])</pre>
for(i in 1:length(unique.d)){
```

returnDcoord 19

returnDcoord

Return D-plane coordinates

# **Description**

Returns the D-plane coordinates given spline parameter values calculated using FindSplineParFNEXTR or FindSplineParFNCOR.

# Usage

```
returnDcoord(par, Gcoords, whichm, sphere.dis = F)
returnDGridcoord(par, gridcoord, Gcoords, whichm, sphere.dis = F)
```

# **Arguments**

par Spline parameter values calculated using either FindSplineParFNEXTR or FindSplineParFNCOR or sdf.heur.Cor or sdf.heur.EXTR.

Goods A d by 2 matrix of G-plane sampling locations.

whichm A vector of length m < d giving the indices of the anchor points in Gcoords.

Must be the same as used to find par.

sphere.dis Is Spherical distance or Euclidean distance used?

gridcoord A K by 2 matrix of any coordinates in the G-plane.

#### Value

returnDcoord returns a d by 2 matrix of coordinates. returnDGridcoord returns a K by 2 matrix of coordinates.

20 Score\_Est

Score\_Est

Score function estimation

#### **Description**

For Score\_Est, the score for the likelihood of a Brown-Resnick or inverted Brown-Resnick model is estimated at each time point. This can then be used to calculate the CLAIC, see examples. Score\_Est\_Smith estimates the score for the likelihood of a Smith or inverted Smith model.

#### Usage

```
Score_Est(Z_Exp, u, coord, lik, type = c("BR,IBR"), sphere.dis = F)
Score_Est_Smith(
    Z_Exp,
    u,
    coord,
    lik,
    type = c("Smith,InvSmith"),
    sphere.dis = F
)
```

#### **Arguments**

Z\_Exp A N by d matrix of exponential random variables.
 u Censoring threshold.
 coord A d by 2 matrix of coordinates.
 lik Optim output from model fitting. See help(nllMSPexp) or help(nllMSPexpSmith).
 type Either Brown-Resnick ("BR") or inverted Brown-Resnick ("IBR") for Score\_Est. Smith or inverted Smith for Score\_Est\_Smith.
 sphere.dis Is Spherical distance or Euclidean distance used?

#### Value

An N by 2 matrix of score estimates for semivariogram parameters  $(\kappa, \lambda)$  (just  $\lambda$  for Score\_Est\_Smith).

```
# Z_Exp: N by d matrix of data on exponential margins
# Gcoords and Dcoords: d by 2 matrix of sampling locations in the G-plane
# and D-plane respectively. See help(returnDcoord) for obtaining Dcoords.
# likG.MSP and likD.MSP: optim outputs from model fitting. See help(nllMSPexp).
#We use data(Aus_Heat) as an example.
##THIS WILL TAKE A LONG TIME TO RUN WITH THE FULL DATASET##

data(Aus_Heat)
Z<-Aus_Heat$Temp.
Gcoords<-Aus_Heat$coords

data(Aus_Heat_Output)</pre>
```

Score\_Est 21

```
sdf<-Aus_Heat_Output$sdf
likG.MSP<-Aus_Heat_Output$likG.MSP
likD.MSP<-Aus_Heat_Output$likD.MSP
Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)
unif<-function(x) rank(x)/(length(x)+1)</pre>
Z U<-Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins
Z_Exp < -qexp(Z_U) #Transform to exponential margins
q <- 0.98 # 98% quantile used as threshold in composite likelihood
u <- quantile(Z_Exp,prob=q)</pre>
s_G_MSP <- Score_Est(Z_Exp,u,coord=Gcoords,lik=likG.MSP,type="BR",sphere.dis=T)</pre>
s_D_MSP <- Score_Est(Z_Exp,u,coord=Dcoords,lik=likD.MSP,type="BR",sphere.dis=T)</pre>
#Estimate variance of score - This is specfic to the Australian summer temperatures data.
# Set block.size. Here we take 90 and 91, corresponding to a regular season
block.sizes \leftarrow c(91,90)
#and a season with a leap year
years <- 1957:2014
k <- length(years)</pre>
temp <- matrix(0,nrow=k,ncol=2)</pre>
temp2 <- matrix(0,nrow=k,ncol=2)</pre>
int <- 0
for(1 in 1:k){
if(years[1]%%4==0) block.size=block.sizes[1] else block.size=block.sizes[2]
 int <- int + block.size</pre>
 temp[1,] <- colSums(s_G_MSP[(int-block.size+1):(int),])</pre>
 temp2[1,] <- colSums(s_D_MSP[(int-block.size+1):(int),])</pre>
}
#Estimate variance of score
varS_G_MSP <- var(temp)</pre>
varS_D_MSP <- var(temp2)</pre>
#Estimate CLAIC
CLAIC_G_MSP <-2*likG.MSP$value+2*sum(diag(varS_G_MSP%*%solve(likG.MSP$hessian)))</pre>
CLAIC_D_MSP <- 2*likD.MSP$value+2*sum(diag(varS_D_MSP%*%solve(likD.MSP$hessian)))
#Estimate CLAIC for Inverted Smith model
#'# likG.IMSP and likD.IMSP: optim outputs from model fitting. See help(nllMSPexp).
likG.IMSP<-Aus_Heat_Output$likG.IMSP
likD.IMSP<-Aus_Heat_Output$likD.IMSP
q <- 0.98 # 98% quantile used as threshold in composite likelihood
u <- quantile(Z_Exp,prob=q)</pre>
s\_G\_IMSP <- Score\_Est\_Smith(Z\_Exp,u,coord=Gcoords,lik=likG.IMSP,type="InvSmith",sphere.dis=T)
```

22 sdf.heur.Cor

```
s_D_IMSP <- Score_Est_Smith(Z_Exp,u,coord=Dcoords,lik=likD.IMSP,type="InvSmith",sphere.dis=T)
#Estimate variance of score - This is specfic to the Australian summer temperatures data.
block.sizes <- c(91,90) #Set block.size # Here we take 90 and 91, corresponding to a regular season
#and a season with a leap year
years <- 1957:2014
k <- length(years)</pre>
temp <- matrix(0,nrow=k,ncol=1)</pre>
temp2 <- matrix(0,nrow=k,ncol=1)</pre>
int <- 0
for(1 in 1:k){
if(years[1]%%4==0) block.size=block.sizes[1] else block.size=block.sizes[2]
 int <- int + block.size</pre>
 temp[1] <- sum(s_G_IMSP[(int-block.size+1):(int)])</pre>
 temp2[1] <- sum(s_D_IMSP[(int-block.size+1):(int)])</pre>
}
#'#Estimate variance of score
varS_G_IMSP <- var(temp)</pre>
varS_D_IMSP <- var(temp2)</pre>
#Estimate CLAIC
\label{local_continuous} $$\operatorname{CLAIC_G_IMSP} <-2*likG.IMSP$value+2*sum(diag(varS_G_IMSP%*%solve(likG.IMSP$hessian)))$$
CLAIC_D_IMSP <- 2*likD.IMSP$value+2*sum(diag(varS_D_IMSP%*%solve(likD.IMSP$hessian)))</pre>
```

sdf.heur.Cor

Heuristic for creating bijective deformations (Correlation)

# **Description**

This algorithm uses FindSplineParFNCOR to create a deformation with m.init inital anchor points. The initial anchor points are the last m.init entries of Full.m.ind. After creating this deformation, the spline values are used as initial parameters in creating a deformation with m.init+1 anchor points. This iterative procedure repeats until a deformation with all Full.m.ind anchor points is created.

#### Usage

```
sdf.heur.Cor(
   m.init,
   Full.m.ind,
   Gcoords,
   emp.cor,
   type = c("F-norm", "Smith"),
   n = 0,
   par = NULL,
   sphere.dis = F
)
```

sdf.heur.Cor 23

#### **Arguments**

m.init Number of inital anchor points. Must have m.init < length(Full.m.ind). Full.m.ind Full vector of indices for anchor points in Gcoords. Gcoords A d by 2 matrix of G-plane coordinates. emp.cor A d by d matrix of pairwise empirical correlation values. "F-norm" for Frobenius norm method using theoretical  $\rho(h_{ij}^*)$  from a Matérn type correlation model. "Smith" for original Smith (1996) method, also using Matérn correlation model. Number of data points used to estimate emp. cor. Only necessary for type=="Smith". n par Initial parameters for first deformation. If not stated, initial parameters are given.

sphere.dis Is Spherical distance or Euclidean distance used?

#### Value

List with three elements:

par Spline parameter values.

value Objective value from final optimisation.

m.ind Vector of indices for full set of anchor points.

```
data("Aus_Heat")
Z<-Aus_Heat$Temp.</pre>
Gcoords<-Aus_Heat$coords
Z_U < -Z
unif<-function(x){rank(x)/(length(x)+1)}</pre>
#Transform to uniform margins
for(i in 1:dim(Z_U)[2]){
  Z_U[,i] < -unif(Z[,i])
}
#Transform to Gaussian margins
Z_N<-qnorm(Z_U)</pre>
#Calculate pairwise empirical correlation
\label{eq:cordinate} emp.cor<-matrix(rep(0,dim(Z_N)[2]^2),nrow=dim(Z_N)[2],ncol=dim(Z_N)[2])
for(i in 1:dim(Z_N)[2]){
  for(j in 1:i){
    \texttt{emp.cor[i,j]} \texttt{<-cor}(Z_N[,i],Z_N[,j])
  }
emp.cor<-emp.cor+t(emp.cor)</pre>
```

24 sdf.heur.EXTR

```
m.init<-6
Full.m.ind<-sample(1:dim(Gcoords)[1],m.init+2)

#Transform to D-plane using Smith (1996) method

##WARNING: This may take a while to run.
sdf<-sdf.heur.Cor(m.init,Full.m.ind,Gcoords,emp.cor,type="Smith",n=dim(Z)[1],sphere.dis=TRUE)

#Plot Dcoords

Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)
plot(Dcoords,main="D-plane",ylab="",xlab="")</pre>
```

sdf.heur.EXTR

Heuristic for creating bijective deformations (Extremal)

# **Description**

This algorithm uses FindSplineParFNEXTR to create a deformation with m.init inital anchor points. The initial anchor points are the last m.init entries of Full.m.ind. After creating this deformation, the spline values are used as initial parameters in creating a deformation with m.init + 1 anchor points. This iterative procedure repeats until a deformation with all Full.m.ind anchor points is created.

# Usage

```
sdf.heur.EXTR(
   m.init,
   Full.m.ind,
   Gcoords,
   emp.chi,
   type = c("CHI_BR", "CHI_q_IBR"),
   q = 0,
   par = NULL,
   sphere.dis = F
)
```

# **Arguments**

m.init	Number of inital anchor points. Must have m.init < length(Full.m.ind).
Full.m.ind	Full vector of indices for anchor points in Gcoords.
Gcoords	A d by 2 matrix of G-plane coordinates.
emp.chi	A d by d matrix of pairwise empirical chi values.
type	"CHI_BR" for theoretical $\chi(h_{ij}^*)$ from a Brown-Resnick model. "CHI_q_IBR" for theoretical $\chi_q(h_{ij}^*)$ from an inverted Brown-Resnick model.
q	Threhold for $\chi_q(h_i^*j)$ . Only needed if type=="CHI_q_IBR".
par	Initial parameters for first deformation. If not stated, initial parameters are given.
sphere.dis	Is Spherical distance or Euclidean distance used?

sdf.heur.EXTR 25

#### Value

```
List with three elements:

par Spline parameter values.

value Objective value from final optimisation.

m.ind Vector of indices for full set of anchor points.
```

```
# Note that the given code will not produce the Australian Summer Temperature deformation
# used in the paper. See Data_Example.R and help(Aus_Heat_Output) for the deformation
# used in the paper. This deformation will use much fewer anchor points as the full
# deformation takes a long time to run.
data("Aus_Heat")
Z<-Aus\_Heat$Temp.
Gcoords<-Aus_Heat$coords
Z_U < -Z
unif<-function(x){rank(x)/(length(x)+1)}</pre>
#Transform to uniform margins
for(i in 1:dim(Z_U)[2]){
  Z_U[,i] < -unif(Z[,i])
}
\label{eq:chi.emp} \begin{split} \text{chi.emp} & -\text{function}(\textbf{U}, \textbf{V}, \textbf{z}) \ \text{sum}((\textbf{U} \!\!> \!\! \textbf{z}) \& (\textbf{V} \!\!> \!\! \textbf{z})) / \text{sum}(\textbf{U} \!\!> \!\! \textbf{z}) \end{split}
q<-0.98
#Calculate pairwise empirical chi
emp.chi<-matrix(rep(0,dim(Z_U)[2]^2),nrow=dim(Z_U)[2],ncol=dim(Z_U)[2])
for(i in 1:dim(Z_U)[2]){
  for(j in 1:i){
    emp.chi[i,j] <-chi.emp(Z_U[,i],Z_U[,j],q) \\
  }
}
emp.chi<-emp.chi+t(emp.chi)</pre>
diag(emp.chi)<-diag(emp.chi)/2</pre>
Full.m.ind<-sample(1:dim(Gcoords)[1],m.init+2)</pre>
\hbox{\tt\#Transform to D-plane using Brown-Resnick theoretical chi function}
##WARNING: This may take a while to run.
sdf<-sdf.heur.EXTR(m.init,Full.m.ind,Gcoords,emp.chi,type="CHI_BR",</pre>
                         par=c(.05,.05,0.2,1.6,rep(0,2*m.init-6)),sphere.dis=TRUE)
#Plot Dcoords
Dcoords<-returnDcoord(sdf$par,Gcoords,sdf$m.ind,sphere.dis=TRUE)</pre>
```

26 Snow\_Precip\_Output

```
plot(Dcoords,main="D-plane",ylab="",xlab="")
```

Snow\_Precip

Snowdonia Precipitation Data

# **Description**

Data consist of winter (DJF) 12-hour average precipitation rate (mm/day) taken from the UKCP18 climate projection. Data is produced on  $2.2 \times 2.2 km^2$  grid-boxes, between the years 1980 and 2000.

# Usage

```
data(Snow_Precip)
```

#### **Format**

A list with 2 elements:

**Pr.** A matrix with 3600 rows and 100 columns of precipitation data.

coords A 100 by 2 matrix of lon-lat coordinates, corresponding to the centroid of each grid-box.

#### References

```
Lowe et al. (2018), Met Office, Exter, UK, (pdf)
```

#### **Examples**

```
data(Snow_Precip)
```

Snow\_Precip\_Output

Snowdonia Precipitation outputs

# Description

Outputs from deformations and model fits for data(Snow\_Precip).

# Usage

```
data(Snow_Precip_Output)
```

#### **Format**

A list with 5 elements:

- **likG.IMSP** Optim output from fitting Inverted Brown-Resnick model to G-plane sampling locations. See nllIMSPexp.
- **likG.MSP** Optim output from fitting Brown-Resnick model to G-plane sampling locations. See n11MSPexp.
- **likD.IMSP** Optim output from fitting Inverted Brown-Resnick model to D-plane sampling locations. See nllIMSPexp.
- **likD.MSP** Optim output from fitting Brown-Resnick model to D-plane sampling locations. See nllMSPexp.
- sdf D-plane transformation spline parameters. See sdf. heur.EXTR.

stat.boot 27

#### References

```
Lowe et al. (2018) Met Office, Exter, UK, (pdf)
```

#### **Examples**

```
data(Snow_Precip_Output)
```

stat.boot

Stationary Bootstrap

#### **Description**

Creates one bootstrap sample of a N by d matrix, X. Here N denotes the number of time points and d the number of sampling locations. The stationary bootstrap (Politis and Romano, 1994) generates a bootstrap sample by repeated sampling of random blocks with expected value block. size until a sample of length N has been created.

# Usage

```
stat.boot(X, mean.block.size)
```

# **Arguments**

Matrix with N rows corresponding to time points and d columns corresponding to spatial locations.

mean.block.size

Expected value of temporal block size.

#### References

```
Politis and Romano (1994), JASA,5(4):303-336, (doi)
```

```
\eqn{N}{} by \eqn{d}{} matrix

data(Aus_Heat)
Z<-Aus_Heat$Temp.

unif<-function(x) rank(x)/(length(x)+1)
Z_U<-Z
for(i in 1:dim(Z_U)[2]) Z_U[,i]<-unif(Z[,i]) # Transform to uniform margins

q<-0.95

ind.triple<-c(1,2,3) ##Denotes indices of triple for which triple-wise chi calculated.

block.mean<-14 #Mean block size for random block choice - Here a fortnight

boot <- stat.boot(Z_U[,ind.triple],block.mean)

#Create one bootstrap estimate of triple-wise chi
chi3.emp(boot[,1],boot[,2],boot[,3],q)</pre>
```

Vterm Vterm

Vterm

Exponent functions

# Description

Exponent function for the Brown-Resnick process and its first- and second-order partial derivatives. For use in likelihoods.

# Usage

```
Vterm(x, y, a)
Vpart(x, y, a)
Vpart2(x, y, a)
```

# **Arguments**

```
x x component. y y component. \sqrt{2\gamma(s_x-s_y)}, \text{ where } s_x \text{ and } s_y \text{ are the sampling locations for the } x \text{ and } y components.
```

# Value

Value

# **Index**

stat.boot, 27

```
Vpart (Vterm), 28
* datasets
    Aus_Heat, 2
                                                    Vpart2 (Vterm), 28
    High_Precip, 11
                                                    Vterm, 28
    Snow_Precip, 26
* output
    Aus_Heat_Output, 2
    High_Precip_Output, 11
    Snow_Precip_Output, 26
Aus_Heat, 2
{\tt Aus\_Heat\_Output, 2}
brnsims, 3
chi3, 4
chi3.emp, 5
chi3q (chi3), 4
FindSplineParFNCOR, 6
{\tt FindSplineParFNEXTR}, 8
gamsv, 9
High_Precip, 11
High\_Precip\_Output, 11
m.reject, 12
makepairs, 13
nl1HT, 14
nllIMSPexp, 15
\verb|nllIMSPexpSmith|, 16
nllMSPexp, 17
\verb|nllMSPexpSmith| (\verb|nllIMSPexpSmith|), 16
returnDcoord, 19
returnDGridcoord (returnDcoord), 19
Score_Est, 20
Score_Est_Smith (Score_Est), 20
sdf.heur.Cor, 22
sdf.heur.EXTR, 24
Snow_Precip, 26
Snow_Precip_Output, 26
```