

# Package ‘sdfEXTREME’

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**Type** Package

**Title** Spatial Deformation for Non-Stationary Extremal Dependence

**Version** 0.1.1

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**Description** Contains functions for creating spatial deformations to account for non-stationarity in spatial data. Both correlation based methods and methods adapted for extremal dependence are included. Accompanies the paper of the same title.

**License** GPL

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Depends** R (>= 2.10)

**Imports** fields,mvtnorm,evd,rmutil,MASS,maps,mapproj,mapdata,numDeriv

**NeedsCompilation** no

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Aus_Heat	<i>Australia Summer Temperatures</i>
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**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)

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Aus_Heat_Output	<i>Australia Summer Temperatures outputs</i>
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**Description**

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

**Usage**

data(Aus\_Heat\_Output)

**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	<i>Simulate non-stationary Brown-Resnick process</i>
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**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**

<code>reps</code>	Number of realisations, $n$ .
<code>locs</code>	A $p$ by 2 matrix of coordinates.
<code>kappa</code>	Value of $\kappa$ . See <code>help(gamsv)</code> .
<code>lambda</code>	Value of $\lambda$ . See <code>help(gamsv)</code> .
<code>tau</code>	A $p$ by $p$ matrix of correlation values.
<code>centre</code>	Vector of length 2 giving coordinates of centre of non-stationarity for semivariogram. If <code>NULL</code> , stationary semivariogram used. See <code>help(gamsv)</code> .

**Value**

$n$  by  $p$  matrix

## 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)))

p<-dim(sim.coords)[1]
tau<-matrix(NA,nrow=p,ncol=p)

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)+
      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.

Sim<-brnsims(reps=10,locs=sim.coords,kappa=kappa,lambda=lambda,centre=centre,tau=tau)
```

---

chi3	<i>Theoretical triple-wise <math>\chi</math> (<math>\chi_q</math>) for a (Inverted) Brown-Resnick process</i>
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---

## 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 $\chi_q(s_i, s_j, s_k)$ .

**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))
```

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.
V	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)
```

---

FindSplineParFNCOR	<i>Find spline parameters (Correlation)</i>
--------------------	---

---

## Description

Find spline parameters for deformation based on correlation methods. Either use the Frobenius 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

<code>par</code>	Parameter values $\phi = (b_1, b_2, \rho, \kappa, \delta_4^{(1)} \dots \delta_m^{(1)}, \delta_4^{(2)} \dots \delta_m^{(2)})$ . If $m < 4$ , $\delta$ parameters are not needed.
<code>Gcoords</code>	A $d$ by 2 matrix of G-plane coordinates.
<code>m.ind</code>	A vector of length $m < d$ giving the indices of the anchor points in <code>Gcoords</code> .
<code>n</code>	Number of data points used to estimate <code>emp.cor</code> . Only necessary for <code>type=="Smith"</code> .
<code>emp.cor</code>	A $d$ by $d$ matrix of pairwise empirical correlation values.
<code>type</code>	"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.
<code>sphere.dis</code>	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.

**Examples**

```
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]){

  Z_U[,i]<-unif(Z[,i])
}
#Transform to Gaussian margins

Z_N<-qnorm(Z_U)

#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)
diag(emp.cor)<-diag(emp.cor)/2

# Set number of anchor points
m<-10
# Sample anchor points
m.ind<-sample(1:dim(Gcoords)[1],m,replace=FALSE)

#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 Frobenius norm of difference between pairwise empirical 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 $\phi = (b_1, b_2, \rho, \kappa, \delta_4^{(1)} \dots \delta_m^{(1)}, \delta_4^{(2)} \dots \delta_m^{(2)})$ . 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	Threshold for $\chi_q(h_{ij}^*)$ . Only needed if <code>type=="CHI_q_IBR"</code> .
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`.

### Examples

```
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]){
```



```

    Z_U[,i]<-unif(Z[,i])
  }

  chi.emp<-function(U,V,z) sum((U>=z)&(V>=z))/sum(U>=z)
  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)
  diag(emp.chi)<-diag(emp.chi)/2

  # Set number of anchor points
  m<-10
  # Sample anchor points
  m.ind<-sample(1:dim(Gcoords)[1],m,replace=FALSE)

  #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,
    Gcoords=Gcoords,m.ind=m.ind,control=list(maxit=2000), emp.dep=emp.chi,
    method = "Nelder-Mead")

  sdf<-optim(fn=FindSplineParFNEXTR,par=sdf$par,sphere.dis=TRUE,type="CHI_BR",
    Gcoords=Gcoords,m.ind=m.ind,
    control=list(maxit=2000), emp.dep=emp.chi,method = "BFGS")
  sdf$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  $s_1$  and  $s_2$ . 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  $\kappa \in (0, 2]$ .

## Usage

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

## Arguments

s1	Vector of length 2 giving coordinates of first location.
s2	Vector of length 2 giving coordinates of second location.
centre	Vector of length 2 giving coordinates of centre of non-stationarity. If NULL, returns stationary semivariogram.
lam	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](#))

## Examples

```
##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)))

p<-dim(sim.coords)[1]
tau<-matrix(NA,nrow=p,ncol=p)

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)+
      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.

Sim<-brnsims(reps=10,locs=sim.coords,kappa=kappa,lambda=lambda,centre=centre,tau=tau)
```

---

High_Precip	<i>Highlands Precipitation Data</i>
-------------	-------------------------------------

---

**Description**

Data consist of winter (DJF) 12-hour average precipitation rate (mm/day) taken from the UKCP18 climate projection. Data is produced on  $2.2 \times 2.2 \text{ km}^2$  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, Exeter, UK, ([pdf](#))

**Examples**

```
data(High_Precip)
```

---

High_Precip_Output	<i>Highlands Precipitation outputs</i>
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---

**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`.

## References

Lowe et al. (2018) Met Office, Exter, UK, ([pdf](#))

## Examples

```
data(High_Precip_Output)
```

---

m.reject	<i>A heuristic for deciding if initial anchor points have enough spread.</i>
----------	--

---

## 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.

## Examples

```
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)
```

makepairs

*Make pairs***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  $u$ .

**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##

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)
```

---

nllHT	<i>Negative log-likelihood for bivariate Heffernan and Tawn (2004) model</i>
-------	--

---

### 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, $Y$ .
par	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](#))

### Examples

```
#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$Temp.
Gcoords<-Aus_Heat$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

p<-dim(Gcoords)[1]
ConExp<-matrix(0,nrow=p,ncol=p)

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),]
```

```

    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="")

```

---

nllIMSPexp	<i>Negative composite censored log-likelihood for the stationary inverted Brown-Resnick process</i>
------------	---

---

## 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.

## 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$Temp.[,1:3]
Gcoords<-Aus_Heat$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)

#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))
dist2<-apply(Zpair[[4]][,3:4],1,function(x){dist[x[1],x[2]]})
unique.d<-as.matrix(unique(dist2))
n.a<-rep(0, dim(unique.d)[1])
for(i in 1:length(unique.d)){
  n.a[i]<-sum(dist2==unique.d[i,1])
}

#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]]),
  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.



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$Temp.[,1:3]
Gcoords<-Aus_Heat$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)

#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))
dist2<-apply(Zpair[[4]][,3:4],1,function(x){dist[x[1],x[2]]})
unique.d<-as.matrix(unique(dist2))
n.a<-rep(0, dim(unique.d)[1])
for(i in 1:length(unique.d)){
  n.a[i]<-sum(dist2==unique.d[i,1])
}

#WARNING- pairwise likelihoods take some time to run
# Inverted Smith process fit to G-plane coordinate system
likG.INV.SMITH<-optim(fn=nllMSPexpSmith,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)
```

## Description

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

## Usage

```
nlMSPexp(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 Brown-Resnick process.

## 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$Temp[,1:3]
Gcoords<-Aus_Heat$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)

#Identify number of non-exceedances per pair
#Speeds up likelihood computation
dist<-rdist(Gcoords+runif(dim(Gcoords)[1]*2,0,1))
dist2<-apply(Zpair[[4]][,3:4],1,function(x){dist[x[1],x[2]]})
unique.d<-as.matrix(unique(dist2))
n.a<-rep(0, dim(unique.d)[1])
for(i in 1:length(unique.d)){
```

```

    n.a[i]<-sum(dist2==unique.d[i,1])
  }

#WARNING- pairwise likelihoods take some time to run
# Brown-Resnick process fit to G-plane coordinate system
likG.MSP<-optim(fn=nllMSPexp,par=c(1.4,500),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 = "Nelder-Mead",hessian=TRUE)

```

---

returnDcoord	<i>Return D-plane coordinates</i>
--------------	-----------------------------------

---

## 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.
Gcoords	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.

---

Score_Est	<i>Score function estimation</i>
-----------	----------------------------------

---

**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).

**Examples**

```
# 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)
```

```

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)
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)

s_G_MSP <- Score_Est(Z_Exp,u,coord=Gcoords,lik=likG.MSP,type="BR",sphere.dis=T)
s_D_MSP <- Score_Est(Z_Exp,u,coord=Dcoords,lik=likD.MSP,type="BR",sphere.dis=T)

#Estimate variance of score - This is specific to the Australian summer temperatures data.
# Set block.size. Here we take 90 and 91, corresponding to a regular season
block.sizes <- c(91,90)
#and a season with a leap year
years <- 1957:2014
k <- length(years)

temp <- matrix(0,nrow=k,ncol=2)
temp2 <- matrix(0,nrow=k,ncol=2)
int <- 0
for(l in 1:k){
  if(years[l]%4==0) block.size=block.sizes[1] else block.size=block.sizes[2]

  int <- int + block.size
  temp[l,] <- colSums(s_G_MSP[(int-block.size+1):(int),])
  temp2[l,] <- colSums(s_D_MSP[(int-block.size+1):(int),])
}

#Estimate variance of score

varS_G_MSP <- var(temp)
varS_D_MSP <- var(temp2)

#Estimate CLAIC
CLAIC_G_MSP <- 2*likG.MSP$value+2*sum(diag(varS_G_MSP%%solve(likG.MSP$hessian)))
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)
s_G_IMSP <- Score_Est_Smith(Z_Exp,u,coord=Gcoords,lik=likG.IMSP,type="InvSmith",sphere.dis=T)

```

```

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 specific 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)

temp <- matrix(0,nrow=k,ncol=1)
temp2 <- matrix(0,nrow=k,ncol=1)
int <- 0
for(l in 1:k){
  if(years[l]%4==0) block.size=block.sizes[1] else block.size=block.sizes[2]

  int <- int + block.size
  temp[l] <- sum(s_G_IMSP[(int-block.size+1):(int)])
  temp2[l] <- sum(s_D_IMSP[(int-block.size+1):(int)])
}

##Estimate variance of score

varS_G_IMSP <- var(temp)
varS_D_IMSP <- var(temp2)

#Estimate CLAIC
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)))

```

---

sdf.heur.Cor

*Heuristic for creating bijective deformations (Correlation)*


---

## Description

This algorithm uses FindSplineParFNCOR to create a deformation with `m.init` initial 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
)

```

## Arguments

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

## Examples

```
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]){

  Z_U[,i]<-unif(Z[,i])
}

#Transform to Gaussian margins

Z_N<-qnorm(Z_U)

#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)
```

```

diag(emp.cor)<-diag(emp.cor)/2

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="")

```

sdf.heur.EXTR

*Heuristic for creating bijective deformations (Extremal)*

### Description

This algorithm uses FindSplineParFNEXTR to create a deformation with `m.init` initial 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

<code>m.init</code>	Number of initial anchor points. Must have <code>m.init &lt; length(Full.m.ind)</code> .
<code>Full.m.ind</code>	Full vector of indices for anchor points in <code>Gcoords</code> .
<code>Gcoords</code>	A <code>d</code> by 2 matrix of G-plane coordinates.
<code>emp.chi</code>	A <code>d</code> by <code>d</code> matrix of pairwise empirical chi values.
<code>type</code>	"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.
<code>q</code>	Threshold for $\chi_q(h_i^*j)$ . Only needed if <code>type=="CHI_q_IBR"</code> .
<code>par</code>	Initial parameters for first deformation. If not stated, initial parameters are given.
<code>sphere.dis</code>	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.

**Examples**

```
# 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)}
```

```
#Transform to uniform margins
for(i in 1:dim(Z_U)[2]){
```

```
  Z_U[,i]<-unif(Z[,i])
}
```

```
chi.emp<-function(U,V,z) sum((U>=z)&(V>=z))/sum(U>=z)
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)
diag(emp.chi)<-diag(emp.chi)/2
```

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

```
#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",
  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)
```

```
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  $2.2 \times 2.2 \text{ km}^2$  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, Exeter, 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 `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`.

## References

Lowe et al. (2018) Met Office, Exter, UK, ([pdf](#))

## Examples

```
data(Snow_Precip_Output)
```

---

stat.boot	<i>Stationary Bootstrap</i>
-----------	-----------------------------

---

## 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

`X` 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](#))

## Examples

```
\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)
```

---

Vterm	<i>Exponent functions</i>
-------	---------------------------

---

**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.

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

**Value**

Value

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