

# Package ‘DiceView’

January 16, 2023

**Title** Methods for Visualization of Computer Experiments Design and Surrogate

**Version** 2.1-0

**Date** 2023-01-16

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**Description** View 2D/3D sections, contour plots, mesh of excursion sets for computer experiments designs, surrogates or test functions.

**Depends** methods, utils, stats, grDevices, graphics

**Imports** DiceDesign, R.cache, geometry, scatterplot3d, parallel

**Suggests** rlikkriging, DiceKriging, DiceEval

**Enhances** rgl, arrangements

**License** GPL-3

**URL** <https://github.com/IRSN/DiceView>

**Repository** CRAN

**RoxygenNote** 7.2.1

**Encoding** UTF-8

**NeedsCompilation** no

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**Date/Publication** 2023-01-16 17:50:02 UTC

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Apply.function	<i>Apply Functions Over Array Margins, using custom vectorization (possibly using parallel)</i>
----------------	---

---

### Description

Emulate parallel apply on a function, from mclapply. Returns a vector or array or list of values obtained by applying a function to margins of an array or matrix.

### Usage

```
Apply.function(
  FUN,
  X,
  MARGIN = 1,
  .combine = c,
  .lapply = parallel::mclapply,
  ...
)
```

### Arguments

FUN	function to apply on X
X	array of input values for FUN
MARGIN	1 indicates to apply on rows (default), 2 on columns
.combine	how to combine results (default using c(.))
.lapply	how to vectorize FUN call (default is parallel::mclapply)
...	optional arguments to FUN.

**Value**

array of values taken by FUN on each row/column of X

**Examples**

```
X = matrix(runif(10),ncol=2);
rowSums(X) == apply(X,1,sum)
apply(X,1,sum) == Apply.function(sum,X)
```

```
X = matrix(runif(10),ncol=1)
rowSums(X) == apply(X,1,sum)
apply(X,1,sum) == Apply.function(sum,X)
```

```
X = matrix(runif(10),ncol=2)
f = function(X) X[1]/X[2]
apply(X,1,f) == Apply.function(f,X)
```

---

are\_in.mesh

*Checks if some points belong to a given mesh*


---

**Description**

Checks if some points belong to a given mesh

**Usage**

```
are_in.mesh(X, mesh)
```

**Arguments**

X	points to check
mesh	mesh identifying the set which X may belong

**Examples**

```
X = matrix(runif(100),ncol=2);
inside = are_in.mesh(X,mesh=geometry::delauayn(matrix(c(0,0,1,1,0,0),ncol=2),output.options =TRUE))
print(inside)
plot(X,col=rgb(1-inside,0,0+inside))
```

---

branin	<i>This is a simple copy of the Branin-Hoo 2-dimensional test function, as provided in DiceKriging package. The Branin-Hoo function is defined here over [0,1] x [0,1], instead of [-5,0] x [10,15] as usual. It has 3 global minima : x1 = c(0.9616520, 0.15); x2 = c(0.1238946, 0.8166644); x3 = c(0.5427730, 0.15)</i>
--------	---

---

**Description**

This is a simple copy of the Branin-Hoo 2-dimensional test function, as provided in DiceKriging package. The Branin-Hoo function is defined here over [0,1] x [0,1], instead of [-5,0] x [10,15] as usual. It has 3 global minima : x1 = c(0.9616520, 0.15); x2 = c(0.1238946, 0.8166644); x3 = c(0.5427730, 0.15)

**Usage**

```
branin(x)
```

**Arguments**

x	a 2-dimensional vector specifying the location where the function is to be evaluated.
---	---

**Value**

A real number equal to the Branin-Hoo function values at x

---

combn.design	<i>Generalize expand.grid() for multi-columns data. Build all combinations of lines from X1 and X2. Each line may hold multiple columns.</i>
--------------	--

---

**Description**

Generalize expand.grid() for multi-columns data. Build all combinations of lines from X1 and X2. Each line may hold multiple columns.

**Usage**

```
combn.design(X1, X2)
```

**Arguments**

X1	variable values, possibly with many columns
X2	variable values, possibly with many columns combn.design(matrix(c(10,20),ncol=1),matrix(c(1,2,3,4,5,6),ncol=2)) combn.design(matrix(c(10,20,30,40),ncol=2),matrix(c(1,2,3,4,5,6),ncol=2))

---

contourview.function *Plot a contour view of a prediction model or function, including design points if available.*

---

### Description

Plot a contour view of a prediction model or function, including design points if available.

### Usage

```
## S3 method for class ``function``
contourview(
  fun,
  vectorized = FALSE,
  dim = NULL,
  center = NULL,
  axis = NULL,
  npoints = 20,
  nlevels = 10,
  col_surf = "blue",
  filled = FALSE,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'matrix'
contourview(
  X,
  y,
  sdy = NULL,
  center = NULL,
  axis = NULL,
  col_points = "red",
  bg_blend = 1,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  title = NULL,
  add = FALSE,
  ...
)
```

```
## S3 method for class 'km'
contourview(
  km_model,
  type = "UK",
  center = NULL,
  axis = NULL,
  npoints = 20,
  nlevels = 10,
  col_points = "red",
  col_surf = "blue",
  filled = FALSE,
  bg_blend = 1,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'Kriging'
contourview(
  Kriging_model,
  center = NULL,
  axis = NULL,
  npoints = 20,
  nlevels = 10,
  col_points = "red",
  col_surf = "blue",
  filled = FALSE,
  bg_blend = 1,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'NuggetKriging'
contourview(
  NuggetKriging_model,
  center = NULL,
  axis = NULL,
  npoints = 20,
```

```
nlevels = 10,  
col_points = "red",  
col_surf = "blue",  
filled = FALSE,  
bg_blend = 1,  
mfrow = NULL,  
Xlab = NULL,  
ylab = NULL,  
Xlim = NULL,  
title = NULL,  
add = FALSE,  
...  
)  
  
## S3 method for class 'NoiseKriging'  
contourview(  
  NoiseKriging_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  nlevels = 10,  
  col_points = "red",  
  col_surf = "blue",  
  filled = FALSE,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  title = NULL,  
  add = FALSE,  
  ...  
)  
  
## S3 method for class 'glm'  
contourview(  
  glm_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  nlevels = 10,  
  col_points = "red",  
  col_surf = "blue",  
  filled = FALSE,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,
```

```

    Xlim = NULL,
    title = NULL,
    add = FALSE,
    ...
)

## S3 method for class 'list'
contourview(
  modelFit_model,
  center = NULL,
  axis = NULL,
  npoints = 20,
  nlevels = 10,
  col_points = "red",
  col_surf = "blue",
  bg_blend = 1,
  filled = FALSE,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

contourview(...)

```

### Arguments

fun	a function or 'predict()'-like function that returns a simple numeric or mean and standard error: list(mean=...,se=...).
vectorized	is fun vectorized?
dim	input variables dimension of the model or function.
center	optional coordinates (as a list or data frame) of the center of the section view if the model's dimension is > 2.
axis	optional matrix of 2-axis combinations to plot, one by row. The value NULL leads to all possible combinations i.e. choose(D, 2).
npoints	an optional number of points to discretize plot of response surface and uncertainties.
nlevels	number of contour levels to display.
col_surf	color for the surface.
filled	use filled.contour
mfrow	an optional list to force par(mfrow = ...) call. The default value NULL is automatically set for compact view.
Xlab	an optional list of string to overload names for X.



<code>ylab</code>	an optional string to overload name for y.
<code>Xlim</code>	an optional list to force x range for all plots. The default value NULL is automatically set to include all design points.
<code>title</code>	an optional overload of main title.
<code>add</code>	to print graphics on an existing window.
<code>...</code>	arguments of the <code>contourview.km</code> , <code>contourview.glm</code> , <code>contourview.Kriging</code> or <code>contourview.function</code> function
<code>X</code>	the matrix of input design.
<code>y</code>	the array of output values.
<code>sd</code>	optional array of output standard error.
<code>col_points</code>	color of points.
<code>bg_blend</code>	an optional factor of alpha (color channel) blending used to plot design points outside from this section.
<code>km_model</code>	an object of class "km".
<code>type</code>	the kriging type to use for model prediction.
<code>Kriging_model</code>	an object of class "Kriging".
<code>NuggetKriging_model</code>	an object of class "Kriging".
<code>NoiseKriging_model</code>	an object of class "Kriging".
<code>glm_model</code>	an object of class "glm".
<code>modelFit_model</code>	an object returned by <code>DiceEval::modelFit</code> .

### Details

If available, experimental points are plotted with fading colors. Points that fall in the specified section (if any) have the color specified `col_points` while points far away from the center have shaded versions of the same color. The amount of fading is determined using the Euclidean distance between the plotted point and center.

### Author(s)

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### See Also

[sectionview.function](#) for a section plot, and [sectionview3d.function](#) for a 2D section plot. [Vectorize.function](#) to wrap as vectorized a non-vectorized function.

[sectionview.matrix](#) for a section plot, and [sectionview3d.matrix](#) for a 2D section plot.

[sectionview.km](#) for a section plot, and [sectionview3d.km](#) for a 2D section plot.

[sectionview.Kriging](#) for a section plot, and [sectionview3d.Kriging](#) for a 2D section plot.

[sectionview.NuggetKriging](#) for a section plot, and [sectionview3d.NuggetKriging](#) for a 2D section plot.

[sectionview.NoiseKriging](#) for a section plot, and [sectionview3d.NoiseKriging](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

### Examples

```
x1 <- rnorm(15)
x2 <- rnorm(15)

y <- x1 + x2 + rnorm(15)
model <- lm(y ~ x1 + x2)

contourview(function(x) sum(x),
             dim=2, Xlim=cbind(range(x1),range(x2)), col='black')
points(x1,x2)

contourview(function(x) {
  x = as.data.frame(x)
  colnames(x) <- names(model$coefficients[-1])
  p = predict.lm(model, newdata=x, se.fit=TRUE)
  list(mean=p$fit, se=p$se.fit)
}, vectorized=TRUE, dim=2, Xlim=cbind(range(x1),range(x2)), add=TRUE)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

contourview(X, y)

if (requireNamespace("DiceKriging")) { library(DiceKriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- km(design = X, response = y, covtype="matern3_2")

contourview(model)

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- Kriging(X = X, y = y, kernel="matern3_2")

contourview(model)

}
```

```

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NuggetKriging(X = X, y = y, kernel="matern3_2")

contourview(model)

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NoiseKriging(X = X, y = y, kernel="matern3_2", noise=rep(5^2,15))

contourview(model)

}

x1 <- rnorm(15)
x2 <- rnorm(15)

y <- x1 + x2^2 + rnorm(15)
model <- glm(y ~ x1 + I(x2^2))

contourview(model)

if (requireNamespace("DiceEval")) { library(DiceEval)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- modelFit(X, y, type = "StepLinear")

contourview(model)

}

## A 2D example - Branin-Hoo function
contourview(branin, dim=2, nlevels=30, col='black')

## Not run:
## a 16-points factorial design, and the corresponding response
d <- 2; n <- 16
design.fact <- expand.grid(seq(0, 1, length = 4), seq(0, 1, length = 4))
design.fact <- data.frame(design.fact); names(design.fact) <- c("x1", "x2")
y <- branin(design.fact); names(y) <- "y"

if (requireNamespace("DiceKriging")) { library(DiceKriging)
## model: km

```

```

model <- DiceKriging::km(design = design.fact, response = y)
contourview(model, nlevels=30)
contourview(branin, dim=2, nlevels=30, col='red', add=TRUE)
}

## model: Kriging
if (requireNamespace("rlikkriging")) { library(rlikkriging)
model <- Kriging(X = as.matrix(design.fact), y = as.matrix(y), kernel="matern3_2")
contourview(model, nlevels=30)
contourview(branin, dim=2, nlevels=30, col='red', add=TRUE)
}

## model: glm
model <- glm(y ~ 1+ x1 + x2 + I(x1^2) + I(x2^2) + x1*x2, data=cbind(y,design.fact))
contourview(model, nlevels=30)
contourview(branin, dim=2, nlevels=30, col='red', add=TRUE)

if (requireNamespace("DiceEval")) { library(DiceEval)
## model: StepLinear
model <- modelFit(design.fact, y, type = "StepLinear")
contourview(model, nlevels=30)
contourview(branin, dim=2, nlevels=30, col='red', add=TRUE)
}

## End(Not run)

```

---

is\_in.mesh

*Checks if some point belongs to a given mesh*


---

## Description

Checks if some point belongs to a given mesh

## Usage

```
is_in.mesh(x, mesh)
```

## Arguments

x	point to check
mesh	mesh identifying the set which X may belong

## Examples

```

is_in.mesh(-0.5, mesh=geometry::deLaunayn(matrix(c(0,1), ncol=1), output.options =TRUE))
is_in.mesh(0.5, mesh=geometry::deLaunayn(matrix(c(0,1), ncol=1), output.options =TRUE))

x =matrix(-.5, ncol=2, nrow=1)
is_in.mesh(x, mesh=geometry::deLaunayn(matrix(c(0,0,1,1,0,0), ncol=2), output.options =TRUE))

```

```
x =matrix(.5,ncol=2,nrow=1)
is_in.mesh(x,mesh=geometry::deilaunayn(matrix(c(0,0,1,1,0,0),ncol=2),output.options =TRUE))
```

---

is\_in.p                      *Test if points are in a hull*

---

### Description

Test if points are in a hull

### Usage

```
is_in.p(x, p, h = NULL)
```

### Arguments

x	points to test
p	points defining the hull
h	hull itself (built from p if given as NULL (default))

### Examples

```
is_in.p(x=-0.5,p=matrix(c(0,1),ncol=1))
is_in.p(x=0.5,p=matrix(c(0,1),ncol=1))
is_in.p(x=matrix(-.5,ncol=2,nrow=1),p=matrix(c(0,0,1,1,0,0),ncol=2))
is_in.p(x=matrix(.25,ncol=2,nrow=1),p=matrix(c(0,0,1,1,0,0),ncol=2))
is_in.p(x=matrix(-.5,ncol=3,nrow=1),p=matrix(c(0,0,0,1,0,0,0,1,0,0,0,1),ncol=3,byrow = TRUE))
is_in.p(x=matrix(.25,ncol=3,nrow=1),p=matrix(c(0,0,0,1,0,0,0,1,0,0,0,1),ncol=3,byrow = TRUE))
```

---

Memoize.function            *Memoize a function*

---

### Description

Before each call of a function, check that the cache holds the results and returns it if available. Otherwise, compute f and cache the result for next evaluations.

### Usage

```
Memoize.function(fun)
```

### Arguments

fun	function to memoize
-----	---------------------

**Value**

a function with same behavior than argument one, but using cache.

**Examples**

```
f=function(n) rnorm(n);
F=Memoize.function(f);
F(5); F(6); F(5)
```

---

 mesh\_exsets

*Search excursion set of nD function, sampled by a mesh*


---

**Description**

Search excursion set of nD function, sampled by a mesh

**Usage**

```
mesh_exsets(
  f,
  vectorized = FALSE,
  threshold,
  sign,
  intervals,
  mesh = "seq",
  mesh.sizes = 11,
  maxerror_f = 1e-09,
  tol = .Machine$double.eps^0.25,
  ex_filter.tri = all,
  ...
)
```

**Arguments**

f	Function to inverse at 'threshold'
vectorized	is f already vectorized ? (default: no)
threshold	target value to inverse
sign	focus at conservative for above (sign=1) or below (sign=-1) the threshold
intervals	bounds to inverse in, each column contains min and max of each dimension
mesh	function or "unif" or "seq" (default) to perform interval partition
mesh.sizes	number of parts for mesh (duplicate for each dimension if using "seq")
maxerror_f	maximal tolerance on f precision
tol	the desired accuracy (convergence tolerance on f arg).
ex_filter.tri	boolean function to validate a geometry::tri as considered in excursion : 'any' or 'all'
...	parameters to forward to mesh_roots(...) call

**Examples**

```

# mesh_exsets(function(x) x, threshold=.51, sign=1, intervals=rbind(0,1),
# maxerror_f=1E-2,tol=1E-2) # for faster testing
# mesh_exsets(function(x) x, threshold=.50000001, sign=1, intervals=rbind(0,1),
# maxerror_f=1E-2,tol=1E-2) # for faster testing
# mesh_exsets(function(x) sum(x), threshold=.51,sign=1, intervals=cbind(rbind(0,1),rbind(0,1)),
# maxerror_f=1E-2,tol=1E-2) # for faster testing
# mesh_exsets(sin, threshold=0, sign="sup", interval=c(pi/2,5*pi/2),
# maxerror_f=1E-2,tol=1E-2) # for faster testing

if (identical(Sys.getenv("NOT_CRAN"), "true")) { # too long for CRAN on Windows

  e = mesh_exsets(function(x) (0.25+x[1])^2+(0.5+x[2])^2 ,
                  threshold =0.25,sign=-1, intervals=matrix(c(-1,1,-1,1),nrow=2),
                  maxerror_f=1E-2,tol=1E-2) # for faster testing

  plot(e$p,xlim=c(-1,1),ylim=c(-1,1));
  apply(e$tri,1,function(tri) polygon(e$p[tri,],col=rgb(.4,.4,.4)))

  if (requireNamespace("rgl")) {
    e = mesh_exsets(function(x) (0.5+x[1])^2+(-0.5+x[2])^2+(0.+x[3])^2,
                    threshold = .25,sign=-1, mesh="unif",
                    intervals=matrix(c(-1,1,-1,1,-1,1),nrow=2),
                    maxerror_f=1E-2,tol=1E-2) # for faster testing

    rgl::plot3d(e$p,xlim=c(-1,1),ylim=c(-1,1),zlim=c(-1,1));
    apply(e$tri,1,function(tri)rgl::lines3d(e$p[tri,]))
  }
}

```

---

 mesh\_roots

*Multi Dimensional Multiple Roots (Zero) Finding, sampled by a mesh*


---

**Description**

Multi Dimensional Multiple Roots (Zero) Finding, sampled by a mesh

**Usage**

```

mesh_roots(
  f,
  vectorized = FALSE,
  intervals,
  mesh = "seq",
  mesh.sizes = 11,
  maxerror_f = 1e-07,
  tol = .Machine$double.eps^0.25,
  ...
)

```

**Arguments**

f	Function (one or more dimensions) to find roots of
vectorized	is f already vectorized ? (default: no)
intervals	bounds to inverse in, each column contains min and max of each dimension
mesh	function or "unif" or "seq" (default) to preform interval partition
mesh.sizes	number of parts for mesh (duplicate for each dimension if using "seq")
maxerror_f	the maximum error on f evaluation (iterates over uniroot to converge).
tol	the desired accuracy (convergence tolerance on f arg).
...	Other args for f

**Value**

matrix of x, so  $f(x)=0$

**Examples**

```

mesh_roots(function(x) x-.51, intervals=rbind(0,1))
mesh_roots(function(x) sum(x)-.51, intervals=cbind(rbind(0,1),rbind(0,1)))
mesh_roots(sin,intervals=c(pi/2,5*pi/2))
mesh_roots(f = function(x) sin(pi*x[1])*sin(pi*x[2]),
           intervals = matrix(c(1/2,5/2,1/2,5/2),nrow=2))

r = mesh_roots(function(x) (0.25+x[1])^2+(0.5+x[2])^2-.25,
              intervals=matrix(c(-1,1,-1,1),nrow=2))
plot(r,xlim=c(-1,1),ylim=c(-1,1))

r = mesh_roots(function(x) (0.5+x[1])^2+(-0.5+x[2])^2+(0.+x[3])^2-.25,
              mesh.sizes = 11,
              intervals=matrix(c(-1,1,-1,1,-1,1),nrow=2))
scatterplot3d::scatterplot3d(r,xlim=c(-1,1),ylim=c(-1,1),zlim=c(-1,1))

mesh_roots(function(x)exp(x)-1,intervals=c(-1,2))
mesh_roots(function(x)exp(1000*x)-1,intervals=c(-1,2))

```

---

min\_dist

*Minimal distance between one point to many points*


---

**Description**

Minimal distance between one point to many points

**Usage**

```
min_dist(x, X, norm = rep(1, ncol(X)))
```



**Arguments**

x	one point
X	matrix of points (same number of columns than x)
norm	normalization vector of distance (same number of columns than x)

**Value**

minimal distance

**Examples**

```
min_dist(runif(3),matrix(runif(30),ncol=3))
```

---

plot2d\_mesh                    *Plot a two dimensional mesh*

---

**Description**

Plot a two dimensional mesh

**Usage**

```
plot2d_mesh(mesh, color = "black", ...)
```

**Arguments**

mesh	2-dimensional mesh to draw
color	color of the mesh
...	optional arguments passed to plot function

**Examples**

```
plot2d_mesh(mesh_exsets(f = function(x) sin(pi*x[1])*sin(pi*x[2]),
                        threshold=0,sign=1, mesh="unif",mesh.size=11,
                        intervals = matrix(c(1/2,5/2,1/2,5/2),nrow=2)))
```

---

plot3d_mesh	<i>Plot a three dimensional mesh</i>
-------------	--------------------------------------

---

**Description**

Plot a three dimensional mesh

**Usage**

```
plot3d_mesh(mesh, engine3d = NULL, color = "black", ...)
```

**Arguments**

mesh	3-dimensional mesh to draw
engine3d	3d framework to use: 'rgl' if installed or 'scatterplot3d' (default)
color	color of the mesh
...	optional arguments passed to plot function

**Examples**

```
if (identical(Sys.getenv("NOT_CRAN"), "true")) { # too long for CRAN on Windows

  plot3d_mesh(mesh_exsets(function(x) (0.5+x[1])^2+(-0.5+x[2])^2+(0.+x[3])^2,
    threshold = .25,sign=-1, mesh="unif",
    maxerror_f=1E-2,tol=1E-2, # faster display
    intervals=matrix(c(-1,1,-1,1,-1,1),nrow=2)),
    engine3d='scatterplot3d')

  if (requireNamespace("rgl")) {
    plot3d_mesh(mesh_exsets(function(x) (0.5+x[1])^2+(-0.5+x[2])^2+(0.+x[3])^2,
      threshold = .25,sign=-1, mesh="unif",
      maxerror_f=1E-2,tol=1E-2, # faster display
      intervals=matrix(c(-1,1,-1,1,-1,1),nrow=2)),engine3d='rgl')
  }
}
```

---

plot_mesh	<i>Plot a one dimensional mesh</i>
-----------	------------------------------------

---

**Description**

Plot a one dimensional mesh

**Usage**

```
plot_mesh(mesh, y = 0, color = "black", ...)
```

**Arguments**

mesh	1-dimensional mesh to draw
y	ordinate value where to draw the mesh
color	color of the mesh
...	optional arguments passed to plot function

**Examples**

```
plot_mesh(mesh_exsets(function(x) x, threshold=.51, sign=1, intervals=rbind(0,1)))
plot_mesh(mesh_exsets(function(x) (x-.5)^2, threshold=.1, sign=-1, intervals=rbind(0,1)))
```

---

points_in.mesh	<i>Extract points of mesh which belong to the mesh triangulation (may not contain all points)</i>
----------------	---

---

**Description**

Extract points of mesh which belong to the mesh triangulation (may not contain all points)

**Usage**

```
points_in.mesh(mesh)
```

**Arguments**

mesh	mesh (list(p,tri,...) from geometry)
------	--------------------------------------

**Value**

points coordinates inside the mesh triangulation

---

points_out.mesh	<i>Extract points of mesh which do not belong to the mesh triangulation (may not contain all points)</i>
-----------------	--

---

**Description**

Extract points of mesh which do not belong to the mesh triangulation (may not contain all points)

**Usage**

```
points_out.mesh(mesh)
```

**Arguments**

mesh	(list(p,tri,...) from geometry)
------	---------------------------------

**Value**

points coordinates outside the mesh triangulation

---

root

*One Dimensional Root (Zero) Finding*

---

**Description**

Search one root with given precision (on y). Iterate over uniroot as long as necessary.

**Usage**

```
root(
  f,
  lower,
  upper,
  maxerror_f = 1e-07,
  f_lower = f(lower, ...),
  f_upper = f(upper, ...),
  tol = .Machine$double.eps^0.25,
  convexity = 0,
  ...
)
```

**Arguments**

<code>f</code>	the function for which the root is sought.
<code>lower</code>	the lower end point of the interval to be searched.
<code>upper</code>	the upper end point of the interval to be searched.
<code>maxerror_f</code>	the maximum error on f evaluation (iterates over uniroot to converge).
<code>f_lower</code>	the same as <code>f(lower)</code> .
<code>f_upper</code>	the same as <code>f(upper)</code> .
<code>tol</code>	the desired accuracy (convergence tolerance on f arg).
<code>convexity</code>	the learned convexity factor of the function, used to reduce the boundaries for uniroot.
<code>...</code>	additional named or unnamed arguments to be passed to f.

**Author(s)**

Yann Richet, IRSN

**Examples**

```
f=function(x) {cat("f");1-exp(x)}; f(root(f,lower=-1,upper=2))
f=function(x) {cat("f");exp(x)-1}; f(root(f,lower=-1,upper=2))

.f = function(x) 1-exp(1*x)
f=function(x) {cat("f");y=.f(x);points(x,y,pch=20,col=rgb(0,0,0,.2));y}
plot(.f,xlim=c(-1,2)); f(root(f,lower=-1,upper=2))

.f = function(x) exp(10*x)-1
f=function(x) {cat("f");y=.f(x);points(x,y,pch=20);y}
plot(.f,xlim=c(-1,2)); f(root(f,lower=-1,upper=2))

.f = function(x) exp(100*x)-1
f=function(x) {cat("f");y=.f(x);points(x,y,pch=20);y}
plot(.f,xlim=c(-1,2)); f(root(f,lower=-1,upper=2))

f=function(x) {cat("f");exp(100*x)-1}; f(root(f,lower=-1,upper=2))
```

---

 roots

*One Dimensional Multiple Roots (Zero) Finding*


---

**Description**

Search multiple roots of 1D function, sampled/splitted by a (1D) mesh

**Usage**

```
roots(
  f,
  interval,
  maxerror_f = 1e-07,
  split = "seq",
  split.size = 11,
  tol = .Machine$double.eps^0.25,
  ...
)
```

**Arguments**

f	Function to find roots
interval	bounds to inverse in
maxerror_f	the maximum error on f evaluation (iterates over uniroot to converge).
split	function or "unif" or "seq" (default) to preform interval partition
split.size	number of parts to perform uniroot inside
tol	the desired accuracy (convergence tolerance on f arg).
...	additional named or unnamed arguments to be passed to f.

**Value**

array of x, so  $f(x)=\text{target}$

**Examples**

```
roots(sin,interval=c(pi/2,5*pi/2))
roots(sin,interval=c(pi/2,1.5*pi/2))
```

```
f=function(x)exp(x)-1;
f(roots(f,interval=c(-1,2)))
```

```
f=function(x)exp(1000*x)-1;
f(roots(f,interval=c(-1,2)))
```

---

sectionview.function *Plot a section view of a prediction model or function, including design points if available.*

---

**Description**

Plot a section view of a prediction model or function, including design points if available.

**Usage**

```
## S3 method for class ``function``
sectionview(
  fun,
  vectorized = FALSE,
  dim = NULL,
  center = NULL,
  axis = NULL,
  npoints = 100,
  col_surf = "blue",
  conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),
  conf_blend = NULL,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  ylim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'matrix'
sectionview(
```

```
X,  
y,  
sdy = NULL,  
center = NULL,  
axis = NULL,  
npoints = 100,  
col_points = "red",  
conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),  
conf_blend = NULL,  
bg_blend = 5,  
mfrow = NULL,  
Xlab = NULL,  
ylab = NULL,  
Xlim = NULL,  
ylim = NULL,  
title = NULL,  
add = FALSE,  
...  
)
```

```
## S3 method for class 'km'  
sectionview(  
  km_model,  
  type = "UK",  
  center = NULL,  
  axis = NULL,  
  npoints = 100,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),  
  conf_blend = NULL,  
  bg_blend = 5,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,  
  ...  
)
```

```
## S3 method for class 'Kriging'  
sectionview(  
  Kriging_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 100,
```

```
col_points = "red",
col_surf = "blue",
conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),
conf_blend = NULL,
bg_blend = 5,
mfrow = NULL,
Xlab = NULL,
ylab = NULL,
Xlim = NULL,
ylim = NULL,
title = NULL,
add = FALSE,
...
)

## S3 method for class 'NuggetKriging'
sectionview(
  NuggetKriging_model,
  center = NULL,
  axis = NULL,
  npoints = 100,
  col_points = "red",
  col_surf = "blue",
  conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),
  conf_blend = NULL,
  bg_blend = 5,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  ylim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'NoiseKriging'
sectionview(
  NoiseKriging_model,
  center = NULL,
  axis = NULL,
  npoints = 100,
  col_points = "red",
  col_surf = "blue",
  conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),
  conf_blend = NULL,
  bg_blend = 5,
  mfrow = NULL,
```



```
    Xlab = NULL,
    ylab = NULL,
    Xlim = NULL,
    ylim = NULL,
    title = NULL,
    add = FALSE,
    ...
)

## S3 method for class 'glm'
sectionview(
  glm_model,
  center = NULL,
  axis = NULL,
  npoints = 100,
  col_points = "red",
  col_surf = "blue",
  conf_lev = c(0.5, 0.8, 0.9, 0.95, 0.99),
  conf_blend = NULL,
  bg_blend = 5,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  ylim = NULL,
  title = NULL,
  add = FALSE,
  ...
)

## S3 method for class 'list'
sectionview(
  modelFit_model,
  center = NULL,
  axis = NULL,
  npoints = 100,
  col_points = "red",
  col_surf = "blue",
  bg_blend = 5,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  ylim = NULL,
  title = NULL,
  add = FALSE,
  ...
)
```

```
sectionview(...)
```

### Arguments

fun	a function or 'predict()'-like function that returns a simple numeric or mean and standard error: list(mean=...,se=...).
vectorized	is fun vectorized?
dim	input variables dimension of the model or function.
center	optional coordinates (as a list or data frame) of the center of the section view if the model's dimension is > 2.
axis	optional matrix of 2-axis combinations to plot, one by row. The value NULL leads to all possible combinations i.e. choose(D, 2).
npoints	an optional number of points to discretize plot of response surface and uncertainties.
col_surf	color for the surface.
conf_lev	an optional list of confidence interval values to display.
conf_blend	an optional factor of alpha (color channel) blending used to plot confidence intervals.
mfrow	an optional list to force par(mfrow = . . .) call. The default value NULL is automatically set for compact view.
Xlab	an optional list of string to overload names for X.
ylab	an optional string to overload name for y.
Xlim	an optional list to force x range for all plots. The default value NULL is automatically set to include all design points.
ylim	an optional list to force y range for all plots.
title	an optional overload of main title.
add	to print graphics on an existing window.
...	arguments of the sectionview.km, sectionview.glm, sectionview.Kriging or sectionview.function function
X	the matrix of input design.
y	the array of output values.
sdv	optional array of output standard error.
col_points	color of points.
bg_blend	an optional factor of alpha (color channel) blending used to plot design points outside from this section.
km_model	an object of class "km".
type	the kriging type to use for model prediction.
Kriging_model	an object of class "Kriging".
NuggetKriging_model	an object of class "Kriging".

NoiseKriging\_model  
                                   an object of class "Kriging".

glm\_model                  an object of class "glm".

modelFit\_model          an object returned by DiceEval::modelFit.

### Details

If available, experimental points are plotted with fading colors. Points that fall in the specified section (if any) have the color specified `col_points` while points far away from the center have shaded versions of the same color. The amount of fading is determined using the Euclidean distance between the plotted point and center.

### Author(s)

Yann Richet, IRSN

### See Also

[sectionview.function](#) for a section plot, and [sectionview3d.function](#) for a 2D section plot. [Vectorize.function](#) to wrap as vectorized a non-vectorized function.

[sectionview.matrix](#) for a section plot, and [sectionview3d.matrix](#) for a 2D section plot.

[sectionview.km](#) for a section plot, and [sectionview3d.km](#) for a 2D section plot.

[sectionview.Kriging](#) for a section plot, and [sectionview3d.Kriging](#) for a 2D section plot.

[sectionview.NuggetKriging](#) for a section plot, and [sectionview3d.NuggetKriging](#) for a 2D section plot.

[sectionview.NoiseKriging](#) for a section plot, and [sectionview3d.NoiseKriging](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

### Examples

```
x1 <- rnorm(15)
x2 <- rnorm(15)

y <- x1 + x2 + rnorm(15)

model <- lm(y ~ x1 + x2)

sectionview(function(x) sum(x),
             dim=2, center=c(0,0), xlim=cbind(range(x1),range(x2)), col='black')

sectionview(function(x) {
  x = as.data.frame(x)
  colnames(x) <- names(model$coefficients[-1])
  p = predict.lm(model, newdata=x, se.fit=TRUE)
  list(mean=p$fit, se=p$se.fit)
}, vectorized=TRUE,
```

```

        dim=2, center=c(0,0), Xlim=cbind(range(x1),range(x2)), add=TRUE)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

sectionview(X,y, center=c(.5,.5))

if (requireNamespace("DiceKriging")) { library(DiceKriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- km(design = X, response = y, covtype="matern3_2")

sectionview(model, center=c(.5,.5))

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- Kriging(X = X, y = y, kernel="matern3_2")

sectionview(model, center=c(.5,.5))

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NuggetKriging(X = X, y = y, kernel="matern3_2")

sectionview(model, center=c(.5,.5))

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NoiseKriging(X = X, y = y, kernel="matern3_2", noise=rep(5^2,15))

sectionview(model, center=c(.5,.5))

}

x1 <- rnorm(15)
x2 <- rnorm(15)

```

```

y <- x1 + x2^2 + rnorm(15)
model <- glm(y ~ x1 + I(x2^2))

sectionview(model, center=c(.5,.5))

if (requireNamespace("DiceEval")) { library(DiceEval)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- modelFit(X, y, type = "StepLinear")

sectionview(model, center=c(.5,.5))

}

## A 2D example - Branin-Hoo function
sectionview(branin, center= c(.5,.5), col='black')

## Not run:
## a 16-points factorial design, and the corresponding response
d <- 2; n <- 16
design.fact <- expand.grid(seq(0, 1, length = 4), seq(0, 1, length = 4))
design.fact <- data.frame(design.fact); names(design.fact) <- c("x1", "x2")
y <- branin(design.fact); names(y) <- "y"

if (requireNamespace("DiceKriging")) { library(DiceKriging)
## model: km
model <- DiceKriging::km(design = design.fact, response = y)
sectionview(model, center= c(.5,.5))
sectionview(branin, center= c(.5,.5), col='red', add=TRUE)
}

if (requireNamespace("rlikkriging")) { library(rlikkriging)
## model: Kriging
model <- Kriging(X = as.matrix(design.fact), y = as.matrix(y), kernel="matern3_2")
sectionview(model, center= c(.5,.5))
sectionview(branin, center= c(.5,.5), col='red', add=TRUE)
}

## model: glm
model <- glm(y ~ 1+ x1 + x2 + I(x1^2) + I(x2^2) + x1*x2, data=cbind(y,design.fact))
sectionview(model, center= c(.5,.5))
sectionview(branin, center= c(.5,.5), col='red', add=TRUE)

if (requireNamespace("DiceEval")) { library(DiceEval)
## model: StepLinear
model <- modelFit(design.fact, y, type = "StepLinear")
sectionview(model, center= c(.5,.5))
sectionview(branin, center= c(.5,.5), col='red', add=TRUE)
}

```

```
## End(Not run)
```

---

```
sectionview3d.function
```

*Plot a contour view of a prediction model or function, including design points if available.*

---

### Description

Plot a contour view of a prediction model or function, including design points if available.

### Usage

```
## S3 method for class ``function``
sectionview3d(
  fun,
  vectorized = FALSE,
  dim = NULL,
  center = NULL,
  axis = NULL,
  npoints = 20,
  col_surf = "blue",
  conf_lev = c(0.95),
  conf_blend = NULL,
  mfrow = NULL,
  Xlab = NULL,
  ylab = NULL,
  Xlim = NULL,
  ylim = NULL,
  title = NULL,
  add = FALSE,
  engine3d = NULL,
  ...
)
```

```
## S3 method for class 'matrix'
sectionview3d(
  X,
  y,
  sdy = NULL,
  center = NULL,
  axis = NULL,
  col_points = "red",
  conf_lev = c(0.95),
  conf_blend = NULL,
  bg_blend = 1,

```

```
mfrow = NULL,  
Xlab = NULL,  
ylab = NULL,  
Xlim = NULL,  
ylim = NULL,  
title = NULL,  
add = FALSE,  
engine3d = NULL,  
...  
)  
  
## S3 method for class 'km'  
sectionview3d(  
  km_model,  
  type = "UK",  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.95),  
  conf_blend = NULL,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,  
  engine3d = NULL,  
  ...  
)  
  
## S3 method for class 'Kriging'  
sectionview3d(  
  Kriging_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.95),  
  conf_blend = NULL,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,
```

```
Xlim = NULL,  
ylim = NULL,  
title = NULL,  
add = FALSE,  
engine3d = NULL,  
...  
)  
  
## S3 method for class 'NuggetKriging'  
sectionview3d(  
  NuggetKriging_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.95),  
  conf_blend = NULL,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,  
  engine3d = NULL,  
  ...  
)  
  
## S3 method for class 'NoiseKriging'  
sectionview3d(  
  NoiseKriging_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.95),  
  conf_blend = NULL,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,
```



```
    engine3d = NULL,  
    ...  
  )  
  
## S3 method for class 'glm'  
sectionview3d(  
  glm_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  conf_lev = c(0.95),  
  conf_blend = NULL,  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,  
  engine3d = NULL,  
  ...  
)  
  
## S3 method for class 'list'  
sectionview3d(  
  modelFit_model,  
  center = NULL,  
  axis = NULL,  
  npoints = 20,  
  col_points = "red",  
  col_surf = "blue",  
  bg_blend = 1,  
  mfrow = NULL,  
  Xlab = NULL,  
  ylab = NULL,  
  Xlim = NULL,  
  ylim = NULL,  
  title = NULL,  
  add = FALSE,  
  engine3d = NULL,  
  ...  
)  
  
sectionview3d(...)
```

**Arguments**

fun	a function or 'predict()-like function that returns a simple numeric or mean and standard error: list(mean=...,se=...).
vectorized	is fun vectorized?
dim	input variables dimension of the model or function.
center	optional coordinates (as a list or data frame) of the center of the section view if the model's dimension is > 2.
axis	optional matrix of 2-axis combinations to plot, one by row. The value NULL leads to all possible combinations i.e. choose(D, 2).
npoints	an optional number of points to discretize plot of response surface and uncertainties.
col_surf	color for the surface.
conf_lev	an optional list of confidence interval values to display.
conf_blend	an optional factor of alpha (color channel) blending used to plot confidence intervals.
mfrow	an optional list to force par(mfrow = ...) call. The default value NULL is automatically set for compact view.
Xlab	an optional list of string to overload names for X.
ylab	an optional string to overload name for y.
Xlim	an optional list to force x range for all plots. The default value NULL is automatically set to include all design points (and their 1-99 percentiles).
ylim	an optional list to force y range for all plots. The default value NULL is automatically set to include all design points (and their 1-99 percentiles).
title	an optional overload of main title.
add	to print graphics on an existing window.
engine3d	3D view package to use. "rgl" if available, otherwise "scatterplot3d" by default.
...	arguments of the sectionview3d.km, sectionview3d.glm, sectionview3d.Kriging or sectionview3d.function function
X	the matrix of input design.
y	the array of output values.
sdv	optional array of output standard error.
col_points	color of points.
bg_blend	an optional factor of alpha (color channel) blending used to plot design points outside from this section.
km_model	an object of class "km".
type	the kriging type to use for model prediction.
Kriging_model	an object of class "Kriging".
NuggetKriging_model	an object of class "Kriging".
NoiseKriging_model	an object of class "Kriging".
glm_model	an object of class "glm".
modelFit_model	an object returned by DiceEval::modelFit.

**Details**

If available, experimental points are plotted with fading colors. Points that fall in the specified section (if any) have the color specified `col_points` while points far away from the center have shaded versions of the same color. The amount of fading is determined using the Euclidean distance between the plotted point and center.

**Author(s)**

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**See Also**

[sectionview.function](#) for a section plot, and [sectionview3d.function](#) for a 2D section plot. [Vectorize.function](#) to wrap as vectorized a non-vectorized function.

[sectionview.matrix](#) for a section plot, and [sectionview3d.matrix](#) for a 2D section plot.

[sectionview.km](#) for a section plot, and [sectionview3d.km](#) for a 2D section plot.

[sectionview.Kriging](#) for a section plot, and [sectionview3d.Kriging](#) for a 2D section plot.

[sectionview.NuggetKriging](#) for a section plot, and [sectionview3d.NuggetKriging](#) for a 2D section plot.

[sectionview.NoiseKriging](#) for a section plot, and [sectionview3d.NoiseKriging](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

[sectionview.glm](#) for a section plot, and [sectionview3d.glm](#) for a 2D section plot.

**Examples**

```
x1 <- rnorm(15)
x2 <- rnorm(15)

y <- x1 + x2 + rnorm(15)
DiceView:::open3d(); DiceView:::plot3d(x1,x2,y)

model <- lm(y ~ x1 + x2)

sectionview3d(function(x) sum(x),
              dim=2, Xlim=cbind(range(x1),range(x2)), add=TRUE, col='black')

sectionview3d(function(x) {
  x = as.data.frame(x)
  colnames(x) <- names(model$coefficients[-1])
  p = predict.lm(model, newdata=x, se.fit=TRUE)
  list(mean=p$fit, se=p$se.fit)
}, vectorized=TRUE, dim=2, Xlim=cbind(range(x1),range(x2)), add=TRUE)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

sectionview3d(X, y)
```

```
if (requireNamespace("DiceKriging")) { library(DiceKriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- km(design = X, response = y, covtype="matern3_2")

sectionview3d(model)

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- Kriging(X = X, y = y, kernel="matern3_2")

sectionview3d(model)

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NuggetKriging(X = X, y = y, kernel="matern3_2")

sectionview3d(model)

}

if (requireNamespace("rlikkriging")) { library(rlikkriging)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin) + 5*rnorm(15)

model <- NoiseKriging(X = X, y = y, kernel="matern3_2", noise=rep(5^2,15))

sectionview3d(model)

}

x1 <- rnorm(15)
x2 <- rnorm(15)

y <- x1 + x2^2 + rnorm(15)
model <- glm(y ~ x1 + I(x2^2))

sectionview3d(model)
```

```

if (requireNamespace("DiceEval")) { library(DiceEval)

X = matrix(runif(15*2),ncol=2)
y = apply(X,1,branin)

model <- modelFit(X, y, type = "StepLinear")

sectionview3d(model)

}

## A 2D example - Branin-Hoo function
sectionview3d(branin, dim=2, col='black')

## Not run:
## a 16-points factorial design, and the corresponding response
d <- 2; n <- 16
design.fact <- expand.grid(seq(0, 1, length = 4), seq(0, 1, length = 4))
design.fact <- data.frame(design.fact); names(design.fact) <- c("x1", "x2")
y <- branin(design.fact); names(y) <- "y"

if (requireNamespace("DiceKriging")) { library(DiceKriging)
## model: km
model <- DiceKriging::km(design = design.fact, response = y)
sectionview3d(model)
sectionview3d(branin, dim=2, col='red', add=TRUE)
}

if (requireNamespace("rlikkriging")) { library(rlikkriging)
## model: Kriging
model <- rlikkriging::Kriging(X = as.matrix(design.fact), y = as.matrix(y), kernel="matern3_2")
sectionview3d(model)
sectionview3d(branin, dim=2, col='red', add=TRUE)
}

## model: glm
model <- glm(y ~ 1+ x1 + x2 + I(x1^2) + I(x2^2) + x1*x2, data=cbind(y,design.fact))
sectionview3d(model)
sectionview3d(branin, dim=2, col='red', add=TRUE)

if (requireNamespace("DiceEval")) { library(DiceEval)
## model: StepLinear
model <- modelFit(design.fact, y, type = "StepLinear")
sectionview3d(model)
sectionview3d(branin, dim=2, col='red', add=TRUE)
}

## End(Not run)

```

**Description**

Vectorize a d-dimensional (input) function, in the same way that `base::Vectorize` for 1-dimensional functions.

**Usage**

```
Vectorize.function(fun, dim, ...)
```

**Arguments**

<code>fun</code>	'dim'-dimensional function to Vectorize
<code>dim</code>	dimension of input arguments of fun
<code>...</code>	optional args to pass to 'Apply.function()', including <code>.combine</code> , <code>.lapply</code> , or optional args passed to 'fun'.

**Value**

a vectorized function (to be called on matrix argument, on each row)

**Examples**

```
f = function(x)x[1]+1; f(1:10); F = Vectorize.function(f,1);  
F(1:10); #F = Vectorize(f); F(1:10);  
  
f2 = function(x)x[1]+x[2]; f2(1:10); F2 = Vectorize.function(f2,2);  
F2(cbind(1:10,11:20));
```

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