Package ‘gMOIP’

February 21, 2024

Type Package

Title Tools for 2D and 3D Plots of Single and Multi-Objective
Linear/Integer Programming Models

Version 1.5.2

URL https://relund.github.io/gMOIP/, https://github.com/relund/gMOIP/

BugReports https://github.com/relund/gMOIP/issues

Description Make 2D and 3D plots of linear programming (LP),
integer linear programming (ILP), or mixed integer linear programming (MILP) models
with up to three objectives. Plots of both the solution and criterion space are possible.
For instance the non-dominated (Pareto) set for bi-objective LP/ILP/MILP programming models
(see vignettes for an overview). The package also contains an function for checking if a point
is inside the convex hull.

License GPL (>= 3.3.2)

Language en-US

Encoding UTF-8

RoxygenNote 7.3.1

Depends R (>= 3.5.0)

Imports ggrepel, geometry, ggplot2, rgl, MASS, Matrix, grDevices,
stats, Rfast, plyr, tidyselect, tidyr, purrr, dplyr,
rlang, png, sp, eaf

Suggests tikzDevice, grid, gridExtra, knitr, rmarkdown, roxygen2,
ggsci, magrittr, scales, pdfpages, testthat (>= 2.1.0),
webshot2

VignetteBuilder knitr

NeedsCompilation no

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Repository CRAN

Date/Publication 2024-02-21 21:30:05 UTC
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addNDSet

Description

Add discrete points to a non-dominated set and classify them into extreme supported, non-extreme supported, non-supported.

Usage

```r
addNDSet(
  pts,
  nDSet = NULL,
  crit = "max",
  keepDom = FALSE,
  dubND = FALSE,
  classify = TRUE
)
```

Arguments

- **pts** A data frame with points to add (a column for each objective).
- **nDSet** A data frame with current non-dominated set (NULL if none yet). Column names of the p objectives must be z1, ..., zp.
- **crit** A max or min vector. If length one assume all objectives are optimized in the same direction.
- **keepDom** Keep dominated points in output.
- **dubND** Duplicated non-dominated points are classified as non-dominated.
- **classify** Non-dominated points are classified into supported extreme (se), supported non-extreme (sne) and unsupported (us).

Value

A data frame with a column for each objective (z columns) and nd (non-dominated). Moreover if classify then columns se, sne, us and cls.

Author(s)

Lars Relund <lars@relund.dk>
Examples

nDSet <- data.frame(z1=c(12,14,16,18), z2=c(18,16,12,4))
pts <- data.frame(z1 = c(18,18,14,15,15), z2=c(2,6,14,14,16))
addNDSet(pts, nDSet, crit = "max")
addNDSet(pts, nDSet, crit = "max", keepDom = TRUE)
addNDSet(pts, nDSet, crit = "min")
addNDSet(c(2,2), nDSet, crit = "max")
addNDSet(c(2,2), nDSet, crit = "min")

nDSet <- data.frame(z1=c(12,14,16,18), z2=c(18,16,12,4), z3 = c(1,7,0,6))
pts <- data.frame(z1=c(12,14,16,18), z2=c(18,16,12,4), z3 = c(2,2,2,6))
crit = c("min", "min", "max")
di <- c(1,1,-1)
li <- c(-1,20)
ini3D(argsPlot3d = list(xlim = li, ylim = li, zlim = li))
plotCones3D(nDSet, direction = di, argsPolygon3d = list(color = "green", alpha = 1),
              drawPoint = FALSE)
plotHull3D(nDSet, addRays = TRUE, direction = di)
plotPoints3D(nDSet, argsPlot3d = list(col = "red"), addText = "coord")
plotPoints3D(pts, addText = "coord")
finalize3D()
addNDSet(pts, nDSet, crit, dubND = FALSE)
addNDSet(pts, nDSet, crit, dubND = TRUE)
addNDSet(pts, nDSet, crit, dubND = TRUE, keepDom = TRUE)
addNDSet(pts, nDSet, crit, dubND = TRUE, keepDom = TRUE, classify = FALSE)

addRays

Add all points on the bounding box hit by the rays.

Description

Add all points on the bounding box hit by the rays.

Usage

addRays(
  pts,
  m = apply(pts, 2, min) - 5,
  M = apply(pts, 2, max) + 5,
  direction = 1
)

Arguments

pts A data frame with all points
m Minimum values of the bounding box.
**binaryPoints**

M
Maximum values of the bounding box.

direction
Ray direction. If i’th entry is positive, consider the i’th column of the pts plus a value greater than or equal zero. If negative, consider the i’th column of the pts minus a value greater than or equal zero.

**Value**

The points merged with the points on the bounding box. The column pt equals 1 if points from pts and zero otherwise.

**Note**

Assume that pts has been checked using `.checkPts()`.

**Examples**

```r
gt <- genNDSet(3,10)[,1:3]
addRays(gt)
addRays(gt, dir = c(1,-1,1))
addRays(gt, dir = c(-1,-1,1), m = c(0,0,0), M = c(100,100,100))
gt <- genSample(5,20)[,1:5]
addRays(gt)
```

---

**binaryPoints**

*Binary (0-1) points in the feasible region (Ax<=b).*

**Description**

Binary (0-1) points in the feasible region (Ax<=b).

**Usage**

`binaryPoints(A, b)`

**Arguments**

- `A`  Constraint matrix.
- `b`  Right hand side.

**Value**

A data frame with all binary points inside the feasible region.

**Note**

Do a simple enumeration of all binary points. Will not work if ncol(A) large.
Author(s)

Lars Relund <lars@relund.dk>.

Examples

```r
A <- matrix(c(3, -2, 1, 2, 4, -2, -3, 2, 1), nc = 3, byrow = TRUE)
b <- c(10, 12, 3)
binaryPoints(A, b)
```

```r
A <- matrix(c(9, 10, 2, 4, -3, 2), ncol = 2, byrow = TRUE)
b <- c(90, 27, 3)
binaryPoints(A, b)
```

---

**classifyNDSet**

*Classify a set of nondominated points*

**Description**

The classification is supported (true/false), extreme (true/false), supported non-extreme (true/false)

**Usage**

```r
classifyNDSet(pts, direction = 1)
```

**Arguments**

- **pts**
  A set of non-dominated points. It is assumed that `ncol(pts)` equals the number of objectives ($p$).

- **direction**
  Ray direction. If i’th entry is positive, consider the i’th column of the `pts` plus a value greater than or equal zero (minimize objective $i$). If negative, consider the i’th column of the `pts` minus a value greater than or equal zero (maximize objective $i$).

**Value**

The ND set with classification columns.

**Note**

It is assumed that `pts` are nondominated.
Examples

```r
classifyNDSet

pts <- matrix(c(0,0,1, 0,1,0, 1,0,0, 0.5,0.2,0.5, 0.25,0.5,0.25), ncol = 3, byrow = TRUE)
ini3D(argsPlot3d = list(xlim = c(min(pts[,1])-2,max(pts[,1])+2),
    ylim = c(min(pts[,2])-2,max(pts[,2])+2),
    zlim = c(min(pts[,3])-2,max(pts[,3])+2)))
plotHull3D(pts, addRays = TRUE, argsPolygon3d = list(alpha = 0.5), useRGLBBox = TRUE)
pts <- classifyNDSet(pts[,1:3])
plotPoints3D(pts$se, col = "red")
plotPoints3D(pts$sne, col = "black")
plotPoints3D(pts$us, col = "blue")
plotCones3D(rectangle = TRUE, argsPolygon3d = list(alpha = 1))
finalize3D()

pts <- matrix(c(0,0,1, 0,1,0, 1,0,0, 0.2,0.1,0.1, 0.1,0.45,0.45), ncol = 3, byrow = TRUE)
di <- -1  # maximize
ini3D(argsPlot3d = list(xlim = c(min(pts[,1])-1,max(pts[,1])+1),
    ylim = c(min(pts[,2])-1,max(pts[,2])+1),
    zlim = c(min(pts[,3])-1,max(pts[,3])+1)))
plotHull3D(pts, addRays = TRUE, argsPolygon3d = list(alpha = 0.5, direction = di),
    addText = "coord")
pts <- classifyNDSet(pts[,1:3], direction = di)
plotPoints3D(pts$se, col = "red")
plotPoints3D(pts$sne, col = "black")
plotPoints3D(pts$us, col = "blue")
plotCones3D(rectangle = TRUE, argsPolygon3d = list(alpha = 1), direction = di)
finalize3D()

pts <- matrix(c(0,0,1, 0,1,0, 1,0,0, 0.5,0.2,0.5, 0.25,0.5,0.25), ncol = 3, byrow = TRUE)
classifyNDSet(pts)

pts <- genNDSet(3,15)
ini3D(argsPlot3d = list(xlim = c(0,max(pts$z1)+2),
    ylim = c(0,max(pts$z2)+2),
    zlim = c(0,max(pts$z3)+2)))
plotHull3D(pts, addRays = TRUE, argsPolygon3d = list(alpha = 0.5))
pts <- classifyNDSet(pts[,1:3])
plotPoints3D(pts$se, col = "red")
plotPoints3D(pts$sne, col = "black")
plotPoints3D(pts$us, col = "blue")
finalize3D()

pts <- genNDSet(3, 15, keepDom = FALSE, argsSphere = list(below = FALSE, factor = 10))
ini3D(argsPlot3d = list(xlim = c(0,max(pts$z1)+2),
    ylim = c(0,max(pts$z2)+2),
    zlim = c(0,max(pts$z3)+2)))
plotHull3D(pts, addRays = TRUE, argsPolygon3d = list(alpha = 0.5))
pts <- classifyNDSet(pts[,1:3])
```
convexHull

Find the convex hull of a set of points.

Description

Find the convex hull of a set of points.

Usage

convexHull(
  pts,
  addRays = FALSE,
  useRGLBox = FALSE,
  direction = 1,
  tol = mean(mean(abs(pts))) * sqrt(.Machine$double.eps) * 2,
  m = apply(pts, 2, min) - 5,
  M = apply(pts, 2, max) + 5
)

Arguments

pts A matrix with a point in each row.
addRays Add the ray defined by direction.
useRGLBox Use the RGL bounding box when add rays.
direction Ray direction. If i'th entry is positive, consider the i'th column of pts plus a value greater than on equal zero (minimize objective $i$). If negative, consider the i'th column of pts minus a value greater than on equal zero (maximize objective $i$).
tol Tolerance on standard deviation if using PCA.
m Minimum values of the bounding box.
M Maximum values of the bounding box.

Value

A list with hull equal a matrix with row indices of the vertices defining each facet in the hull and pts equal the input points (and dummy points) and columns: pt, true if a point in the original input; false if a dummy point (a point on a ray). vtx, TRUE if a vertex in the hull.
cornerPoints

## Examples

### 1D
```r
pts<-matrix(c(1,2,3), ncol = 1, byrow = TRUE)
dimFace(pts) # a line
convexHull(pts)
convexHull(pts, addRays = TRUE)
```

### 2D
```r
pts<-matrix(c(1,1, 2,2), ncol = 2, byrow = TRUE)
dimFace(pts) # a line
convexHull(pts)
plotHull2D(pts, drawPoints = TRUE)
convexHull(pts, addRays = TRUE)
plotHull2D(pts, addRays = TRUE, drawPoints = TRUE)
pts<-matrix(c(1,1, 2,2, 0,1), ncol = 2, byrow = TRUE)
dimFace(pts) # a polygon
convexHull(pts)
plotHull2D(pts, drawPoints = TRUE)
convexHull(pts, addRays = TRUE, direction = c(-1,1))
plotHull2D(pts, addRays = TRUE, direction = c(-1,1), addText = "coord")
```

### 3D
```r
pts<-matrix(c(1,1,1), ncol = 3, byrow = TRUE)
dimFace(pts) # a point
convexHull(pts)
pts<-matrix(c(0,0,0,1,1,1,2,2,2,3,3,3), ncol = 3, byrow = TRUE)
dimFace(pts) # a line
convexHull(pts)
pts<-matrix(c(0,0,0,1,1,0,2,2,0,2,0,2), ncol = 3, byrow = TRUE)
dimFace(pts) # a polygon
convexHull(pts)
convexHull(pts, addRays = TRUE)
pts<-matrix(c(1,0,0,1,1,1,2,2,1,1,1,1), ncol = 3, byrow = TRUE)
dimFace(pts) # a polygon
convexHull(pts)
pts<-matrix(c(1,1,1,2,2,1,1,1,1,1,1,2), ncol = 3, byrow = TRUE)
dimFace(pts) # a polytope (polyhedron)
convexHull(pts)
ini3D(argsPlot3d = list(xlim = c(0,3), ylim = c(0,3), zlim = c(0,3)))
pts<-matrix(c(1,1,1,2,2,1,1,1,1,1,1,2), ncol = 3, byrow = TRUE)
plotPoints3D(pts)
plotHull3D(pts, argsPolygon3d = list(color = "red"))
convexHull(pts)
plotHull3D(pts, addRays = TRUE)
convexHull(pts, addRays = TRUE)
finalize3D()
```

cornerPoints  

*Calculate the corner points for the polytope Ax<=b.*
cornerPointsCont

Description
Calculate the corner points for the polytope Ax\leq b assuming all variables are continuous.

Usage
cornerPointsCont(A, b, nonneg = rep(TRUE, ncol(A)))
**criterionPoints**

**Arguments**

- **A**  
  Constraint matrix.
- **b**  
  Right hand side.
- **nonneg**  
  A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.

**Value**

A data frame with a corner point in each row.

**Author(s)**

Lars Relund <lars@relund.dk>

---

**criterionPoints**  
*Calculate the criterion points of a set of points and ranges to find the set of non-dominated points (Pareto points) and classify them into extreme supported, non-extreme supported, non-supported.*

**Description**

Calculate the criterion points of a set of points and ranges to find the set of non-dominated points (Pareto points) and classify them into extreme supported, non-extreme supported, non-supported.

**Usage**

`criterionPoints(pts, obj, crit, labels = "coord")`

**Arguments**

- **pts**  
  A data frame with a column for each variable in the solution space (can also be a `rangePoints`).
- **obj**  
  A p x n matrix(one row for each criterion).
- **crit**  
  Either max or min.
- **labels**  
  If NULL or "n" don’t add any labels (empty string). If equals coord, labels are the solution space coordinates. Otherwise number all points from one based on the solution space points.

**Value**

A data frame with columns x1, ..., xn, z1, ..., zp, lbl (label), nD (non-dominated), ext (extreme), nonExt (non-extreme).

**Author(s)**

Lars Relund <lars@relund.dk>
Examples

A <- matrix(c(3, -2, 1, 2, 4, -2, -3, 2, 1), nc = 3, byrow = TRUE)
b <- c(10, 12, 3)
pts <- IntegerPoints(A, b)
obj <- matrix(c(1, -3, 1, -1, 1, -1), byrow = TRUE, ncol = 3)
criterionPoints(pts, obj, crit = "max", labels = "numb")

---

**df2String**

Convert each row to a string.

**Description**

Convert each row to a string.

**Usage**

df2String(df, round = 2)

**Arguments**

- **df**: Data frame.
- **round**: How many digits to round.

**Value**

A vector of strings.

---

**dimFace**

Return the dimension of the convex hull of a set of points.

**Description**

Return the dimension of the convex hull of a set of points.

**Usage**

dimFace(pts, dim = NULL)

**Arguments**

- **pts**: A matrix/data frame/vector that can be converted to a matrix with a row for each point.
- **dim**: The dimension of the points, i.e. assume that column 1-dim specify the points. If NULL assume that the dimension are the number of columns.
**finalize3D**

Finalize the RGL window.

**Description**

Finalize the RGL window.

**Usage**

`finalize3D(...)`

**Value**

The dimension of the object.

**Examples**

```r
## In 1D
pts <- matrix(c(3), ncol = 1, byrow = TRUE)
dimFace(pts)
pts <- matrix(c(1,3,4), ncol = 1, byrow = TRUE)
dimFace(pts)

## In 2D
pts <- matrix(c(3,3,6,3,6,3,6,3,6,6,3,6), ncol = 2, byrow = TRUE)
dimFace(pts)
pts <- matrix(c(1,1,2,3,3,3,3,3,3,3,3,3), ncol = 2, byrow = TRUE)
dimFace(pts)
pts <- matrix(c(0,0), ncol = 2, byrow = TRUE)
dimFace(pts)

## In 3D
pts <- c(3,3,3,6,3,3,6,3,6,3,6,6,3)
dimFace(pts, dim = 3)
pts <- matrix( c(1,1,1), ncol = 3, byrow = TRUE)
dimFace(pts)
pts <- matrix( c(1,1,1,2,2,2,2,2,2,2,2,2), ncol = 3, byrow = TRUE)
dimFace(pts)
pts <- matrix(c(2,2,2,2,2,2,2,2,2,2,2,2), ncol=3, byrow= TRUE)
dimFace(pts)
pts <- matrix(c(0,0,0,0,1,1,0,2,2,0,5,0,6,1), ncol = 3, byrow = TRUE)
dimFace(pts)
pts <- matrix(c(0,0,0,0,1,1,0,2,2,0,2,1,1,1), ncol = 3, byrow = TRUE)
dimFace(pts)

## In 4D
pts <- matrix(c(2,2,2,2,2,2,2,2,2,2,2,2,3,4,1,2,3,4), ncol=4, byrow= TRUE)
dimFace(pts)
```
Arguments

Further arguments passed on the RGL plotting functions. This must be done as lists. Currently the following arguments are supported:

- `argsAxes3d`: A list of arguments for `rgl::axes3d`.
- `argsTitle3d`: A list of arguments for `rgl::title3d`.

Value

The RGL object (using `rgl::highlevel()`).

Examples

```r
ini3D()
pts<-matrix(c(1,1,1,5,5,5), ncol = 3, byrow = TRUE)
plotPoints3D(pts)
finalize3D()

ini3D()
pts<-matrix(c(1,1,1,5,5,5), ncol = 3, byrow = TRUE)
plotPoints3D(pts)
finalize3D(argsAxes3d = list(edges = "bbox"))
```

---

**genNDSet**

*Generate a sample of nondominated points.*

Description

Generate a sample of nondominated points.

Usage

```r
genNDSet(
p, 
n, 
range = c(1, 100), 
random = FALSE, 
sphere = TRUE, 
planes = FALSE, 
box = FALSE, 
keepDom = FALSE, 
crit = "min", 
dubND = FALSE, 
classify = FALSE, 
... 
)
```
Arguments

- **p**  Dimension of the points.
- **n**  Number nondominated points generated.
- **range**  The range of the points in each dimension (a vector or matrix with p rows).
- **random**  Random sampling.
- **sphere**  Generate points on a sphere.
- **planes**  Generate points between two planes.
- **box**  Generate points in boxes.
- **keepDom**  Keep dominated points also.
- **crit**  Criteria used (a vector of min/max).
- **dubND**  Should duplicated non-dominated points be considered as non-dominated.
- **classify**  Non-dominated points are classified into supported extreme (se), supported non-extreme (sne) and unsupported (us)
- ...  Further arguments passed on to genSample.

Value

A data frame with p+1 columns (last one indicate if dominated or not).

Examples

```r
## Random
range <- matrix(c(1,100, 50, 100, 10, 50), ncol = 2, byrow = TRUE)
pts <- genNDSet(3, 5, range = range, random = TRUE, keepDom = TRUE)
head(pts)
Rfast::colMinsMaxs(as.matrix(pts[, 1:3]))
iini3D(FALSE, argsPlot3d = list(xlim = c(min(pts[,1])-2,max(pts[,1])+10),
ylim = c(min(pts[,2])-2,max(pts[,2])+10),
zlim = c(min(pts[,3])-2,max(pts[,3])+10))
plotPoints3D(pts[,1:3])
plotPoints3D(pts$nd[,1:3], argsPlot3d = list(col = "red", size = 10))
plotCones3D(pts$nd[,1:3], argsPolygon3d = list(alpha = 1))
final3D()

## Between planes
range <- matrix(c(1,10000, 1,10000), ncol = 2, byrow = TRUE)
pts <- genNDSet(2, 50, range = range, planes = TRUE, classify = TRUE)
head(pts)
Rfast::colMinsMaxs(as.matrix(pts[, 1:2]))
plot(pts[, 1:2])

range <- matrix(c(1,100, 50,100, 10, 50), ncol = 2, byrow = TRUE)
center <- rowMeans(range)
planeU <- c(rep(1, 3), -1.2*sum(rowMeans(range)))
planeL <- c(rep(1, 3), -0.8*sum(rowMeans(range)))
```
genSample

Generate a sample of points in dimension $p$.

**Description**

Generate a sample of points in dimension $p$. 

```r
pts <- genNDSet(3, 50, range = range, planes = TRUE, keepDom = TRUE, classify = TRUE,
  argsPlanes = list(center = center, planeU = planeU, planeL = planeL))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts[, 1:3]))
ini3D(FALSE, argsPlot3d = list(xlim = c(min(pts[,1])-2, max(pts[,1])+10),
  ylim = c(min(pts[,2])-2, max(pts[,2])+10),
  zlim = c(min(pts[,3])-2, max(pts[,3])+10),
  box = TRUE, axes = TRUE))
plotPoints3D(pts[,1:3])
plotPoints3D(pts[pts$nd,1:3], argsPlot3d = list(col = "red", size = 10))
rgl::planes3d(planeL[1], planeL[2], planeL[3], planeL[4], alpha = 0.5)
rgl::planes3d(planeU[1], planeU[2], planeU[3], planeU[4], alpha = 0.5)
finalize3D()

## On a sphere
ini3D()
range <- c(1,100)
cent <- rep(range[1] + (range[2]-range[1])/2, 3)
pts <- genNDSet(3, 20, range = range, sphere = TRUE, keepDom = TRUE,
  argsSphere = list(center = cent))
rgl::spheres3d(cent, radius=49.5, color = "grey100", alpha=0.1)
plotPoints3D(pts)
plotPoints3D(pts[pts$nd,], argsPlot3d = list(col = "red", size = 10))
rgl::planes3d(cent[1],cent[2],cent[3],-sum(cent^2), alpha = 0.5, col = "red")
finalize3D()
ini3D()
cent <- c(100,100,100)
r <- 75
planeC <- c(cent+r/3)
planeC <- c(planeC, -sum(planeC^2))
pts <- genNDSet(3, 20, keepDom = TRUE,
  argsSphere = list(center = cent, radius = r, below = FALSE, plane = planeC, factor = 6))
rgl::spheres3d(cent, radius=r, color = "grey100", alpha=0.1)
plotPoints3D(pts)
plotPoints3D(pts[pts$nd,], argsPlot3d = list(col = "red", size = 10))
rgl::planes3d(planeC[1],planeC[2],planeC[3],planeC[4], alpha = 0.5, col = "red")
finalize3D()
```
Usage

genSample(
  p,
  n,
  range = c(1, 100),
  random = FALSE,
  sphere = TRUE,
  planes = FALSE,
  box = FALSE,
  ...
)

Arguments

p  Dimension of the points.
n  Number of samples generated.
range  The range of the points in each dimension (a vector or matrix with p rows).
random  Random sampling.
sphere  Generate points on a sphere.
planes  Generate points between two planes.
box  Generate points in boxes.
...  Further arguments passed on to the method for generating points. This must be done as lists (see examples). Currently the following arguments are supported:

• argsPlanes: A list of arguments for generating points between planes and in the cube defined by the range:
  – center: A point between the planes (default rowMeans(range)).
  – planeU: The upper plane (default c(rep(1, p), -1.2*sum(center))).
  – planeL: The lower plane (default c(rep(1, p), -0.8*sum(center))).

• argsSphere: A list of arguments for generating points on a sphere:
  – radius: The radius of the sphere.
  – center: The center of the sphere.
  – plane: The plane used.
  – below: Either true (generate points below the plane), false (generate points above the plane) or NULL (generated on the whole sphere).
  – factor: If using a plane. Then the factor to multiply n with, so generate enough points below/above the plane.
  – closeToPlane: If TRUE only return points close to the plane.

• argsBox: A list of arguments for generating points inside boxes:
  – intervals: Number of intervals to split the length of the range into. That is, each range is divided into intervals (sub)intervals and only the lowest/highest subrange is used.
  – cor: How to correlate indices. If 'idxAlt' then alternate the intervals (high/low) for each dimension. For instance if p = 3 and the first dimension is in the high interval range then the second will be in the low
interval range and third in the high interval range again. If \( \text{idxRand} \) then choose the low/high interval range for each dimension based on \( \text{prHigh} \). If \( \text{idxSplit} \) then select \( \lfloor p/2 \rfloor : \lceil p/2 \rceil \) dimensions for the high interval range and the other for the low interval range.

- \( \text{prHigh} \): Probability for choosing the high interval range in each dimension.

**Details**

Note having ranges with different length when using the sphere method, doesn’t make sense. The best option is properly to use a center and radius here. Moreover, as for higher \( p \) you may have to use a larger radius than half of the desired interval range.

**Value**

A matrix with \( p \) columns.

**Examples**

```r
### Using random
## p = 2
range <- matrix(c(1,100, 50,100), ncol = 2, byrow = TRUE )
pts <- genSample(2, 1000, range = range, random = TRUE)
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts)

## p = 3
range <- matrix(c(1,100, 50,100, 10,50), ncol = 2, byrow = TRUE )
ini3D()
pts <- genSample(3, 1000, range = range, random = TRUE)
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plotPoints3D(pts)
finalize3D()

## other p
p <- 10
range <- c(1,100)
pts <- genSample(p, 1000, range = range, random = TRUE)
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))

### Using planes
## p = 2
range <- matrix(c(1,100, 50,100), ncol = 2, byrow = TRUE )
center <- rowMeans(range)
planesL <- c(rep(1, 2), -1.5*sum(rowMeans(range)))
planesU <- c(rep(1, 2), -0.7*sum(rowMeans(range)))
```
```r
pts <- genSample(2, 1000, range = range, planes = TRUE,
                 argsPlanes = list(center = center, planeU = planeU, planeL = planeL))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts)

## p = 3
range <- matrix(c(1,100, 50,100, 10, 50), ncol = 2, byrow = TRUE)
center <- rowMeans(range)
planeU <- c(rep(1, 3), -1.2*sum(rowMeans(range)))
planeL <- c(rep(1, 3), -0.6*sum(rowMeans(range)))
pts <- genSample(3, 1000, range = range, planes = TRUE,
                 argsPlanes = list(center = center, planeU = planeU, planeL = planeL))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
in3D(argsPlot3d = list(box = TRUE, axes = TRUE))
plotPoints3D(pts)
rgl::planes3d(planeL[1], planeL[2], planeL[3], planeL[4], alpha = 0.5)
rgl::planes3d(planeU[1], planeU[2], planeU[3], planeU[4], alpha = 0.5)
finalize3D()

### Using sphere
## p = 2
range <- matrix(c(1,100, 50,100, 10, 50), ncol = 2, byrow = TRUE)
cent <- rep(range[1] + (range[2]-range[1])/2, 2)
pts <- genSample(2, 1000, range = range)
dim(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts, asp=1)
abline(sum(cent^2)/cent[1], -cent[2]/cent[1])

cent <- c(100,100)
r <- 75
planeC <- c(cent+r/3)
planeC <- c(planeC, -sum(planeC^2))
pts <- genSample(2, 100,
                 argsSphere = list(center = cent, radius = r, below = FALSE, plane = planeC, factor = 6))
dim(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts, asp=1)
abline(-planeC[3]/planeC[1], -planeC[2]/planeC[1])

pts <- genSample(2, 100, argsSphere = list(center = cent, radius = r, below = NULL))
dim(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts, asp=1)

## p = 3
ini3D()
```

range <- c(1,100)
cent <- rep(range[1] + (range[2]-range[1])/2, 3)
pts <- genSample(3, 1000, range = range)
dim(pts)
Rfast::colMinsMaxs(as.matrix(pts))
grl::spheres3d(cent, radius=49.5, color = "grey100", alpha=0.1)
plotPoints3D(pts)
grl::planes3d(cent[1],cent[2],cent[3],-sum(cent^2), alpha = 0.5, col = "red")
finalize3D()

ini3D()
cent <- c(100,100,100)
r <- 75
planeC <- c(cent+r/3)
planeC <- c(planeC, -sum(planeC^2))
pts <- genSample(3, 100,
  argsSphere = list(center = cent, radius = r, below = FALSE, plane = planeC, factor = 6))
grl::spheres3d(cent, radius=r, color = "grey100", alpha=0.1)
plotPoints3D(pts)
grl::planes3d(planeC[1],planeC[2],planeC[3],planeC[4], alpha = 0.5, col = "red")
finalize3D()

ini3D()
pts <- genSample(3, 10000, argsSphere = list(center = cent, radius = r, below = NULL))
Rfast::colMinsMaxs(as.matrix(pts))
grl::spheres3d(cent, radius=r, color = "grey100", alpha=0.1)
plotPoints3D(pts)
finalize3D()

## Other p
p <- 10
cent <- rep(0,p)
r <- 100
pts <- genSample(p, 100000, argsSphere = list(center = cent, radius = r, below = NULL))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
apply(pts,1, function(x){sqrt(sum((x-cent)^2))}) # test should be approx. equal to radius

## Using box
## p = 2
range <- matrix(c(1,100, 50,100), ncol = 2, byrow = TRUE )
pts <- genSample(2, 1000, range = range, box = TRUE, argsBox = list(cor = "idxAlt"))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plot(pts)

pts <- genSample(2, 1000, range = range, box = TRUE, argsBox = list(cor = "idxRand"))
plot(pts)
getTexture

plot(pts)
pts <- genSample(2, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxRand", prHigh = c(0.1,0.6)))
points(pts, pch = 3, col = "red")
pts <- genSample(2, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxRand", prHigh = c(0,0)))
points(pts, pch = 4, col = "blue")
pts <- genSample(2, 1000, range = range, box = TRUE, argsBox = list(cor = "idxSplit"))
plot(pts)

## p = 3
range <- matrix(c(1,100, 1,200, 1,50), ncol = 2, byrow = TRUE )
ini3D(argsPlot3d = list(box = TRUE, axes = TRUE))
pts <- genSample(3, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxAlt"))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
plotPoints3D(pts)
finalize3D()
ini3D(argsPlot3d = list(box = TRUE, axes = TRUE))
pts <- genSample(3, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxAlt", intervals = 6))
plotPoints3D(pts)
finalize3D()
ini3D(argsPlot3d = list(box = TRUE, axes = TRUE))
pts <- genSample(3, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxRand"))
plotPoints3D(pts)
pts <- genSample(3, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxRand", prHigh = c(0.1,0.6,0.1)))
plotPoints3D(pts, argsPlot3d = list(col="red"))
finalize3D()
ini3D(argsPlot3d = list(box = TRUE, axes = TRUE))
pts <- genSample(3, 1000, range = range, box = TRUE, 
                argsBox = list(cor = "idxSplit"))
plotPoints3D(pts)
finalize3D()

## other p
p <- 10
range <- c(1,100)
pts <- genSample(p, 1000, range = range, box = TRUE, argsBox = list(cor = "idxSplit"))
head(pts)
Rfast::colMinsMaxs(as.matrix(pts))
**Description**

Save a point symbol as a temporary file.

**Usage**

```r
getAddress(pch = 16, cex = 10, ...)
```

**Arguments**

- `pch`: Point number/symbol.
- `cex`: Point size
- `...`: Further arguments passed to `plot`.

**Value**

The file name.

**Examples**

```r
# Pch shapes
generateRPointShapes<-function(){
  oldPar<-par()
  par(font=2, mar=c(0.5,0,0,0))
  y=rev(c(rep(1,6),rep(2,5), rep(3,5), rep(4,5), rep(5,5)))
  x=c(rep(1:5,5),6)
  plot(x, y, pch = 0:25, cex=1.5, ylim=c(1.5,5.5), xlim=c(1,6.5),
       axes=FALSE, xlab="", ylab="", bg="blue")
  text(x, y, labels=0:25, pos=3)
  par(mar=oldPar$mar,font=oldPar$font)
}
generateRPointShapes()

getAddress()
```

---

**gMOIPTheme**

*The ggplot theme for the package*

**Description**

The ggplot theme for the package

**Usage**

```r
gMOIPTheme(...)
```
## hullSegment

### Arguments

... Further arguments parsed to `ggplot2::theme()`.

### Value

The theme object.

### Examples

```r
pts <- matrix(c(1,1), ncol = 2, byrow = TRUE)
plotHull2D(pts)
pts1 <- matrix(c(2,2, 3,3), ncol = 2, byrow = TRUE)
pts2 <- matrix(c(1,1, 2,2, 0,1), ncol = 2, byrow = TRUE)
ggplot2::ggplot() +
  plotHull2D(pts2, drawPoints = TRUE, addText = "coord", drawPlot = FALSE) +
  plotHull2D(pts1, drawPoints = TRUE, drawPlot = FALSE) +
  gMOIPTheme() +
  gggplot2::xlab(expression(x[1])) +
  gggplot2::ylab(expression(x[2]))
```

### Description

Find segments (lines) of a face.

### Usage

```r
hullSegment(
  vertices,
  hull = geometry::convhulln(vertices),
  tol = mean(mean(abs(vertices))) * sqrt(.Machine$double.eps)
)
```

### Arguments

- **vertices**: A \(m \times p\) array of vertices of the convex hull, as used by `geometry::convhulln()`.
- **hull**: Tessellation (or triangulation) generated by `geometry::convhulln()` If hull is left empty or not supplied, then it will be generated.
- **tol**: Tolerance on the tests for inclusion in the convex hull. You can think of \(tol\) as the distance a point may possibly lie outside the hull, and still be perceived as on the surface of the hull. Because of numerical slop nothing can ever be done exactly here. I might guess a semi-intelligent value of \(tol\) to be \(tol = 1.e-13*\text{mean(abs(\text{vertices}()))}\)

In higher dimensions, the numerical issues of floating point arithmetic will probably suggest a larger value of \(tol\).
inHull

Efficient test for points inside a convex hull in p dimensions.

Description
Efficient test for points inside a convex hull in p dimensions.

Usage

inHull(
  pts,
  vertices,
  hull = NULL,
  tol = mean(mean(abs(as.matrix(vertices)))) * sqrt(.Machine$double.eps)
)

Arguments

- **pts**: A \(n \times p\) array to test, \(n\) data points, in dimension \(p\). If you have many points to test, it is most efficient to call this function once with the entire set.
- **vertices**: A \(m \times p\) array of vertices of the convex hull. May contain redundant (non-vertex) points.
- **hull**: Tessellation (or triangulation) generated by `convhulln` (only works if the dimension of the hull is \(p\)). If hull is NULL, then it will be generated.
- **tol**: Tolerance on the tests for inclusion in the convex hull. You can think of tol as the difference a point value may be different from the values of the hull, and still be perceived as on the surface of the hull. Because of numerical slop nothing can ever be done exactly here. In higher dimensions, the numerical issues of floating point arithmetic will probably suggest a larger value of tol. tol is not used if the dimension of the hull is larger than one and not equal \(p\).

Value
An integer vector of length \(n\) with values 1 (inside hull), -1 (outside hull) or 0 (on hull to precision indicated by tol).

Note
Some of the code are inspired by the Matlab code by John D'Errico and how to find a point inside a hull. If the dimension of the hull is below \(p\) then PCA may be used to check (a warning will be given).
Author(s)

Lars Relund <lars@relund.dk>

Examples

## In 1D
vertices <- matrix(4, ncol = 1)
pt <- matrix(c(2,4), ncol = 1, byrow = TRUE)
inHull(pt, vertices)

## In 2D
vertices <- matrix(c(2,4), ncol = 2)
pt <- matrix(c(2,4, 1,1), ncol = 2, byrow = TRUE)
inHull(pt, vertices)

## In 3D
vertices <- matrix(c(2,2,2), ncol = 3, byrow = TRUE)
pt <- matrix(c(1,1, 1,1, 1,0, 1,0, 1,0,1), ncol = 3, byrow = TRUE)
inHull(pt, vertices)
ini3D

Initialize the RGL window.

Usage

ini3D(new = TRUE, clear = FALSE, ...)

Arguments

new A new window is opened (otherwise the current is cleared).
clear Clear the current RGL window.
... Further arguments passed on the the RGL plotting functions. This must be done
as lists. Currently the following arguments are supported:
  • argsPlot3d: A list of arguments for rgl::plot3d.
  • argsAspect3d: A list of arguments for rgl::aspect3d.

Value

NULL (invisible).

Examples

ini3D()
pts<-matrix(c(1,1,1,5,5,5), ncol = 3, byrow = TRUE)
plotPoints3D(pts)
finalize3D()

lim <- c(-1, 7)
ini3D(argsPlot3d = list(xlim = lim, ylim = lim, zlim = lim))
plotPoints3D(pts)
finalize3D()
integerPoints

Integer points in the feasible region (Ax<=b).

Description

Integer points in the feasible region (Ax<=b).

Usage

integerPoints(A, b, nonneg = rep(TRUE, ncol(A)))

Arguments

A  Constraint matrix.
b  Right hand side.
nonneg A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.

Value

A data frame with all integer points inside the feasible region.

Note

Do a simple enumeration of all integer points between min and max values found using the continuous polytope.

Author(s)

Lars Relund <lars@relund.dk>.

Examples

A <- matrix( c(3,-2, 1, 2, 4,-2,-3, 2, 1), nc = 3, byrow = TRUE)
b <- c(10, 12, 3)
integerPoints(A, b)

A <- matrix(c(9, 10, 2, 4, -3, 2), ncol = 2, byrow = TRUE)
b <- c(90, 27, 3)
integerPoints(A, b)
loadView

Help function to load the view angle for the RGL 3D plot from a file or matrix

Description

Help function to load the view angle for the RGL 3D plot from a file or matrix

Usage

loadView(
  fname = "view.RData",
  v = NULL,
  clear = TRUE,
  close = FALSE,
  zoom = 1,
  ...
)

Arguments

fname The file name of the view.
v The view matrix.
clear Call rgl::clear3d().
close Call rgl::close3d().
zoom Zoom level.
... Additional parameters passed to rgl::view3d().

Author(s)

Lars Relund <lars@relund.dk>

Examples

view <- matrix( c(-0.412063330411911, -0.228006735444069, 0.882166087627411, 0,
                  0.910147845745087, -0.0574885793030262, 0.410274744033813, 0, -0.042830865830183,
                  0.97196090221405, 0.231208890676498, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix( c(3, 2, 5, 2, 1, 1, 1, 3, 5, 2, 4), nc = 3, byrow = TRUE)
b <- c(55, 26, 30, 57)
obj <- c(20, 10, 15)
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord")

# Try to modify the angle in the RGL window
saveView(print = TRUE) # get the view angle to insert into R code
mergeLists  

Merge two lists to one

Description
Merge two lists to one

Usage
mergeLists(a, b)

Arguments
- \(a\)  
First list.
- \(b\)  
Second list.

plotCones2D  

Plot a cone defined by a point in 2D.

Description
The cones are defined as the point plus/minus rays of R2.

Usage
plotCones2D(

  pts,
  drawPoint = TRUE,
  drawLines = TRUE,
  drawPolygons = TRUE,
  direction = 1,
  rectangle = FALSE,
  drawPlot = TRUE,
  m = apply(pts, 2, min) - 5,
  M = apply(pts, 2, max) + 5,
  ...
)

Arguments
- \(pts\)  
A matrix with a point in each row.
- \(drawPoint\)  
Draw the points defining the cone.
- \(drawLines\)  
Draw lines of the cone.
- \(drawPolygons\)  
Draw polygons of the cone.
direction Ray direction. If i'th entry is positive, consider the i'th column of pts plus a value greater than on equal zero (minimize objective i$\$). If negative, consider the i'th column of pts minus a value greater than on equal zero (maximize objective i$\$).

rectangle Draw the cone as a rectangle.

drawPlot Draw the ggplot. Set to FALSE if you want to combine hulls in a single plot.

m Minimum values of the bounding box.

M Maximum values of the bounding box.

... Further arguments passed to plotHull2D

Value

A ggplot object

Examples

library(ggplot2)
plotCones2D(c(4,4), drawLines = FALSE, drawPoint = TRUE,
argsGeom_point = list(col = "red", size = 10),
argsGeom_polygon = list(alpha = 0.5), rectangle = TRUE)
plotCones2D(c(1,1), rectangle = FALSE)
plotCones2D(matrix(c(3,3,2,2), ncol = 2, byrow = TRUE))

## The Danish flag
lst <- list(argsGeom_polygon = list(alpha = 0.85, fill = "red"),
drawPlot = FALSE, drawPoint = FALSE, drawLines = FALSE)
p1 <- do.call(plotCones2D, args = c(list(c(2,4), direction = 1), lst))
p2 <- do.call(plotCones2D, args = c(list(c(1,2), direction = -1), lst))
p3 <- do.call(plotCones2D, args = c(list(c(2,2), direction = c(1,-1)), lst))
p4 <- do.call(plotCones2D, args = c(list(c(1,4), direction = c(-1,1)), lst))
ggplot() + p1 + p2 + p3 + p4 + theme_void()
plotCones3D

  rectangle = FALSE,
  useRGLBBBox = TRUE,
...
)

Arguments

pts A matrix with a point in each row.
drawPoint Draw the points defining the cone.
drawLines Draw lines of the cone.
drawPolygons Draw polygons of the cone.
direction Ray direction. If i’th entry is positive, consider the i’th column of pts plus a value greater than zero (minimize objective $i$). If negative, consider the i’th column of pts minus a value greater than zero (maximize objective $i$).
rectangle Draw the cone as a rectangle.
useRGLBBBox Use the RGL bounding box as ray limits for the cone.
... Further arguments passed on to the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:
  • argsPlot3d: A list of arguments for rgl::plot3d.
  • argsSegments3d: A list of arguments for rgl::segments3d.
  • argsPolygon3d: A list of arguments for rgl::polygon3d.

Value

Object ids (invisible).

Examples

ini3D(argsPlot3d = list(xlim = c(0,6), ylim = c(0,6), zlim = c(0,6)))
plotCones3D(c(4,4,4), drawLines = FALSE, drawPoint = TRUE,
  argsPlot3d = list(col = "red", size = 10),
  argsPolygon3d = list(alpha = 1), rectangle = TRUE)
plotCones3D(c(1,1,1), rectangle = FALSE)
plotCones3D(matrix(c(3,3,3,2,2,2), ncol = 3, byrow = TRUE))
finalize3D()

ini3D(argsPlot3d = list(xlim = c(0,6), ylim = c(0,6), zlim = c(0,6)))
plotCones3D(c(4,4,4), direction = 1)
plotCones3D(c(2,2,2), direction = -1)
plotCones3D(c(4,2,2), direction = c(1,-1,-1))
ids <- plotCones3D(c(2,2,4), direction = c(-1,-1,1))
finalize3D()
# pop3d(id = ids) # remove last cone
plotCriterion2D  

Create a plot of the criterion space of a bi-objective problem

Description
Create a plot of the criterion space of a bi-objective problem

Usage
plotCriterion2D(
  A,
  b,
  obj,
  type = rep("c", ncol(A)),
  nonneg = rep(TRUE, ncol(A)),
  crit = "max",
  addTriangles = FALSE,
  addHull = TRUE,
  plotFeasible = TRUE,
  latex = FALSE,
  labels = NULL
)

Arguments
A  The constraint matrix.
b  Right hand side.
obj A p x n matrix(one row for each criterion).
type A character vector of same length as number of variables. If entry k is ’i’ variable k must be integer and if ’c’ continuous.
nonneg A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.
crit Either max or min (only used if add the iso-profit line).
addTriangles Add search triangles defined by the non-dominated extreme points.
addHull Add the convex hull and the rays.
plotFeasible If True then plot the criterion points/slices.
latex If true make latex math labels for TikZ.
labels If NULL don’t add any labels. If ’n’ no labels but show the points. If equal coord add coordinates to the points. Otherwise number all points from one.

Value
The ggplot object.
Note

Currently only points are checked for dominance. That is, for MILP models some nondominated points may in fact be dominated by a segment.

Author(s)

Lars Relund <lars@relund.dk>

Examples

```r
### Set up 2D plot
# Function for plotting the solution and criterion space in one plot (two variables)
plotBiObj2D <- function(A, b, obj,
  type = rep("c", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = FALSE,
  labels = "numb",
  addTriangles = TRUE,
  addHull = TRUE)
{
  p1 <- plotPolytope(A, b, type = type, crit = crit, faces = faces, plotFaces = plotFaces,
    plotFeasible = plotFeasible, plotOptimum = plotOptimum, labels = labels)
  p2 <- plotCriterion2D(A, b, obj, type = type, crit = crit, addTriangles = addTriangles,
    addHull = addHull, plotFeasible = plotFeasible, labels = labels)
  gridExtra::grid.arrange(p1, p2, nrow = 1)
}

### Bi-objective problem with two variables
A <- matrix(c(-3,2,2,4,9,10), ncol = 2, byrow = TRUE)
b <- c(3,27,90)
## LP model
obj <- matrix(
  c(7, -10, # first criterion
    -10, -10), # second criterion
  nrow = 2)
plotBiObj2D(A, b, obj, addTriangles = FALSE)

## ILP models with different criteria (maximize)
obj <- matrix(c(3, -1, -2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
obj <- matrix(c(-7, -1, -5, 5), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
obj <- matrix(c(-1, -1, 2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
```

```r
# Bi-objective problem with two variables
A <- matrix(c(-3,2,2,4,9,10), ncol = 2, byrow = TRUE)
b <- c(3,27,90)
## LP model
obj <- matrix(
  c(7, -10, # first criterion
    -10, -10), # second criterion
  nrow = 2)
plotBiObj2D(A, b, obj, addTriangles = FALSE)

## ILP models with different criteria (maximize)
obj <- matrix(c(3, -1, -2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
obj <- matrix(c(-7, -1, -5, 5), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
obj <- matrix(c(-1, -1, 2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)))
```
## ILP models with different criteria (minimize)

```r
obj <- matrix(c(7, -10, -10, -10), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)), crit = "min")
obj <- matrix(c(3, -1, -2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)), crit = "min")
obj <- matrix(c(-7, -1, -5, 5), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)), crit = "min")
obj <- matrix(c(-1, -1, 2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = rep("i", ncol(A)), crit = "min")
```

# More examples

## MILP model (x1 integer) with different criteria (maximize)

```r
obj <- matrix(c(7, -10, -10, -10), nrow = 2)
plotBiObj2D(A, b, obj, type = c("i", "c"))
obj <- matrix(c(3, -1, -2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = c("i", "c"))
obj <- matrix(c(-7, -1, -5, 5), nrow = 2)
plotBiObj2D(A, b, obj, type = c("i", "c"))
obj <- matrix(c(-1, -1, 2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = c("i", "c"))
```

## MILP model (x2 integer) with different criteria (minimize)

```r
obj <- matrix(c(7, -10, -10, -10), nrow = 2)
plotBiObj2D(A, b, obj, type = c("c", "i"), crit = "min")
obj <- matrix(c(3, -1, -2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = c("c", "i"), crit = "min")
obj <- matrix(c(-7, -1, -5, 5), nrow = 2)
plotBiObj2D(A, b, obj, type = c("c", "i"), crit = "min")
obj <- matrix(c(-1, -1, 2, 2), nrow = 2)
plotBiObj2D(A, b, obj, type = c("c", "i"), crit = "min")
```

### Set up 3D plot

```r
# Function for plotting the solution and criterion space in one plot (three variables)
plotBiObj3D <- function(A, b, obj,
                     type = rep("c", ncol(A)),
                     crit = "max",
                     faces = rep("c", ncol(A)),
                     plotFaces = TRUE,
                     plotFeasible = TRUE,
                     plotOptimum = TRUE,
                     labels = "numb",
                     addTriangles = TRUE,
                     addHull = TRUE)
{
  plotPolytope(A, b, type = type, crit = crit, faces = faces, plotFaces = plotFaces,
               plotFeasible = plotFeasible, plotOptimum = plotOptimum, labels = labels)
  plotCriterion2D(A, b, obj, type = type, crit = crit, addTriangles = addTriangles,
                  addHull = addHull, plotFeasible = plotFeasible, labels = labels)
}
```
### Bi-objective problem with three variables

```r
loadView <- function(fname = "view.RData", v = NULL) {
  if (!is.null(v)) {
    rgl::view3d(userMatrix = v)
  } else {
    if (file.exists(fname)) {
      load(fname)
      rgl::view3d(userMatrix = view)
    } else {
      warning(paste0("Can't load view in file ", fname, "!"))
    }
  }
}
#
# Ex
view <- matrix(c(-0.452365815639496, -0.446501553058624, 0.77201122045517, 0, 0.886364221572876, 
                 -0.320795893669128, 0.333835482597351, 0, 0.0986008867621422, 0.835299551486969, 
                 0.540881276130676, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
Ab <- matrix(c(1, 1, 2, 5, 
              2, -1, 0, 3, 
              -1, 2, 1, 3, 
              0, -3, 5, 2 
), nc = 4, byrow = TRUE)
A <- Ab[,1:3]
b <- Ab[,4]
obj <- matrix(c(1, -6, 3, -4, 1, 6), nrow = 2)
# LP model
plotBiObj3D(A, b, obj, crit = "min", addTriangles = FALSE)
# ILP model
plotBiObj3D(A, b, obj, type = c("i","i","i"), crit = "min")
# MILP model
plotBiObj3D(A, b, obj, type = c("c","i","i"), crit = "min")
plotBiObj3D(A, b, obj, type = c("i","c","i"), crit = "min")
plotBiObj3D(A, b, obj, type = c("i","i","c"), crit = "min")
plotBiObj3D(A, b, obj, type = c("c","c","i"), crit = "min")

# Ex
view <- matrix(c(0.976349174976349, -0.202332556247711, 0.77201122045517, 0, 0.886364221572876, 
                 -0.320795893669128, 0.333835482597351, 0, 0.0986008867621422, 0.835299551486969, 
                 0.540881276130676, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix(c(-1, 1, 0, 
              1, 4, 0, 
              -1, 0, 3, 
              2, 1, 0, 
              -1, 3, 2, 
              0, -3, 1 
), nc = 4)
```

```r
```
2, 1, 0,
3, -4, 0,
0, 0, 4
}, nc = 3, byrow = TRUE)
b <- c(5, 45, 24, 10)
obj <- matrix(c(1, -6, 3, -4, 1, 6), nrow = 2)
# LP model
plotBiObj3D(A, b, obj, crit = "min", addTriangles = FALSE, labels = "coord")
# ILP model
plotBiObj3D(A, b, obj, type = c("i","i","i"))
# MILP model
plotBiObj3D(A, b, obj, type = c("c","i","i"))
plotBiObj3D(A, b, obj, type = c("i","c","i"), plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("i","i","c"), plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("c","i","c"), plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("c","c","i"))

## Ex
view <- matrix(c(-0.812462985515594, -0.029454167932272, 0.582268416881561, 0, 0.579295456409454,
                 -0.153386667370796, 0.800555109977722, 0, 0.065732568502461, 0.987727105617523,
                 0.14168381690979, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix(c(1, 1, 1,
              3, 0, 1
), nc = 3, byrow = TRUE)
b <- c(10, 24)
obj <- matrix(c(1, -6, 3, -4, 1, 6), nrow = 2)
# LP model
plotBiObj3D(A, b, obj, crit = "min", addTriangles = FALSE, labels = "coord")
# ILP model
plotBiObj3D(A, b, obj, type = c("i","i","i"), crit = "min", labels = "n")
# MILP model
plotBiObj3D(A, b, obj, type = c("c","i","i"), crit = "min")
plotBiObj3D(A, b, obj, type = c("i","c","i"), crit = "min")
plotBiObj3D(A, b, obj, type = c("i","i","c"), crit = "min")
plotBiObj3D(A, b, obj, type = c("i","c","c"), crit = "min")
plotBiObj3D(A, b, obj, type = c("c","i","c"), crit = "min", plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("c","c","i"), crit = "min", plotFaces = FALSE)

## Ex
view <- matrix(c(-0.412063338411911, -0.228006735444069, 0.882166087627411, 0, 0.910147845745087,
                 -0.0574885793030262, 0.410274744833813, 0, -0.042830865830183, 0.97196090221405,
                 0.231208890676498, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix( c(3, 2, 5,
2, 1, 1,
1, 1, 3,
5, 2, 4), nc = 3, byrow = TRUE)
b <- c(55, 26, 30, 57)
obj <- matrix(c(1, -6, 3, -4, 1, -1), nrow = 2)

# LP model
plotBiObj3D(A, b, obj, crit = "min", addTriangles = FALSE, labels = "coord")

# ILP model
plotBiObj3D(A, b, obj, type = c("i","i","i"), crit = "min", labels = "n")

# MILP model
plotBiObj3D(A, b, obj, type = c("c","i","i"), crit = "min", labels = "n")
plotBiObj3D(A, b, obj, type = c("i","c","i"), crit = "min", labels = "n", plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("i","i","c"), crit = "min", labels = "n")
plotBiObj3D(A, b, obj, type = c("c","i","c"), crit = "min", labels = "n", plotFaces = FALSE)
plotBiObj3D(A, b, obj, type = c("c","c","i"), crit = "min", labels = "n")

---

plotHull2D

Plot the convex hull of a set of points in 2D.

Description

Plot the convex hull of a set of points in 2D.

Usage

plotHull2D( 
   pts,
   drawPoints = FALSE,
   drawLines = TRUE,
   drawPolygons = TRUE,
   addText = FALSE,
   addRays = FALSE,
   direction = 1,
   drawPlot = TRUE,
   drawBBoxHull = FALSE,
   m = apply(pts, 2, min) - 5,
   M = apply(pts, 2, max) + 5,
   ... 
)
Arguments

- **pts**: A matrix with a point in each row.
- **drawPoints**: Draw the points.
- **drawLines**: Draw lines of the facets.
- **drawPolygons**: Fill the hull.
- **addText**: Add text to the points. Currently coord (coordinates), rownames (rownames) and both supported or a vector with text.
- **addRays**: Add the ray defined by direction.
- **direction**: Ray direction. If i'th entry is positive, consider the i'th column of pts plus a value greater than on equal zero (minimize objective i$\$). If negative, consider the i'th column of pts minus a value greater than on equal zero (maximize objective i$\$).
- **drawPlot**: Draw the ggplot. Set to FALSE if you want to combine hulls in a single plot.
- **drawBBoxHull**: If addRays then draw the hull areas hitting the bounding box also.
- **m**: Minimum values of the bounding box.
- **M**: Maximum values of the bounding box.
- **...**: Further arguments passed on the the ggplot plotting functions. This must be done as lists. Currently the following arguments are supported:
  - **argsGeom_point**: A list of arguments for `ggplot2::geom_point`.
  - **argsGeom_path**: A list of arguments for `ggplot2::geom_path`.
  - **argsGeom_polygon**: A list of arguments for `ggplot2::geom_polygon`.
  - **argsGeom_label**: A list of arguments for `ggplot2::geom_label`.

Value

The ggplot object if drawPlot = TRUE; otherwise, a list of ggplot components.

Examples

```r
library(ggplot2)
pts <- matrix(c(1,1), ncol = 2, byrow = TRUE)
pHull2D(pts)
pts1 <- matrix(c(2,2, 3,3), ncol = 2, byrow = TRUE)
pHull2D(pts1, drawPoints = TRUE)
pHull2D(pts1, drawPoints = TRUE, addRays = TRUE, addText = "coord")
pHull2D(pts1, drawPoints = TRUE, addRays = TRUE, addText = "coord", drawBBoxHull = TRUE)
pHull2D(pts1, drawPoints = TRUE, addRays = TRUE, direction = -1, addText = "coord")
pts2 <- matrix(c(1,1, 2,2, 0,1), ncol = 2, byrow = TRUE)
pHull2D(pts2, drawPoints = TRUE, addText = "coord")
pHull2D(pts2, drawPoints = TRUE, addRays = TRUE, addText = "coord")
pHull2D(pts2, drawPoints = TRUE, addRays = TRUE, direction = -1, addText = "coord")
## Combine hulls
ggplot() +
pHull2D(pts2, drawPoints = TRUE, addText = "coord", drawPlot = FALSE) +
pHull2D(pts1, drawPoints = TRUE, drawPlot = FALSE) +
gMOIPTheme() +
```

# Plotting an LP
A <- matrix(c(-3,2,2,4,9,10), ncol = 2, byrow = TRUE)
b <- c(3,27,90)
obj <- c(7.75, 10)
pts3 <- cornerPoints(A, b)
plotHull2D(pts3, drawPoints = TRUE, addText = "coord", argsGeom_polygon = list(fill = "red"))

plotHull3D

## Description
Plot the convex hull of a set of points in 3D.

## Usage

plotHull3D(
  pts,
  drawPoints = FALSE,
  drawLines = TRUE,
  drawPolygons = TRUE,
  addText = FALSE,
  addRays = FALSE,
  useRGLBBox = TRUE,
  direction = 1,
  drawBBoxHull = TRUE,
  ...
)

## Arguments

- **pts**: A matrix with a point in each row.
- **drawPoints**: Draw the points.
- **drawLines**: Draw lines of the facets.
- **drawPolygons**: Fill the facets.
- **addText**: Add text to the points. Currently coord (coordinates), rownames (rownames) and both supported or a vector with text.
- **addRays**: Add the ray defined by direction.
- **useRGLBBox**: Use the RGL bounding box when add rays.
- **direction**: Ray direction. If i'th entry is positive, consider the i'th column of pts plus a value greater than or equal zero (minimize objective $i$). If negative, consider the i'th column of pts minus a value greater than or equal zero (maximize objective $i$).
If addRays then draw the hull areas hitting the bounding box also.

Further arguments passed on the the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:

- `argsPlot3d`: A list of arguments for `rgl::plot3d`.
- `argsSegments3d`: A list of arguments for `rgl::segments3d`.
- `argsPolygon3d`: A list of arguments for `rgl::polygon3d`.
- `argsShade3d`: A list of arguments for `rgl::shade3d`.
- `argsText3d`: A list of arguments for `rgl::text3d`.

Value

A list with hull, pts classified and object ids (invisible).

Examples

```r
ini3D()
pts <- matrix(c(0,0,0), ncol = 3, byrow = TRUE)
plotHull3D(pts) # a point
pts <- matrix(c(1,1,1,2,2,3,3), ncol = 3, byrow = TRUE)
plotHull3D(pts, drawPoints = TRUE) # a line
pts <- matrix(c(1,0,1,1,1,2,3,1,1,3,3), ncol = 3, byrow = TRUE)
plotHull3D(pts, drawLines = FALSE, argsPolygon3d = list(alpha=0.6)) # a polygon
lst <- plotHull3D(pts, argsPolygon3d = list(alpha=0.9), argsSegments3d = list(color="red"))
finalize3D()
# pop3d(id = lst$ids) # remove last hull

## Using addRays
pts <- data.frame(x = c(1,3), y = c(1,3), z = c(1,3))
in3D(argsPlot3d = list(xlim = c(0,max(pts$x)+10),
     ylim = c(0,max(pts$y)+10),
     zlim = c(0,max(pts$z)+10))
plotHull3D(pts, drawPoints = TRUE, addRays = TRUE, drawBBoxHull = FALSE)
plotHull3D(c(4,4,4), drawPoints = TRUE, addRays = TRUE)
finalize3D()

pts <- data.frame(x = c(4,2.5,1), y = c(1,2.5,4), z = c(1,2.5,4))
in3D(argsPlot3d = list(xlim = c(0,max(pts$x)+10),
     ylim = c(0,max(pts$y)+10),
     zlim = c(0,max(pts$z)+10))
plotHull3D(pts, drawPoints = TRUE, addRays = TRUE)
finalize3D()

pts <- matrix(c(4,8,0,
               0,4,8,
               8,4,0,
               4,8,0,
               4,0,8,
               8,0,4),
              ncol=3, byrow = TRUE)
```
plotLines2D

Plot the lines of a linear mathematical program (Ax = b)

Description

Plot the lines of a linear mathematical program (Ax = b)
plotMTeX3D

Usage

plotMTeX3D(tex, edge, line = 0, at = NULL, pos = NA, ...)
**plotNDSet2D**  
Create a plot of a discrete non-dominated set.

**Description**  
Create a plot of a discrete non-dominated set.

**Usage**  
plotNDSet2D(
  points,
  crit,
  addTriangles = FALSE,
  addHull = TRUE,
  latex = FALSE,
  labels = NULL
)

**Arguments**  
- **points**: Data frame with non-dominated points.
- **crit**: Either max or min (only used if add the iso-profit line). A vector is currently not supported.
- **addTriangles**: Add search triangles defined by the non-dominated extreme points.
- **addHull**: Add the convex hull and the rays.
- **latex**: If true make latex math labels for TikZ.
- **labels**: If NULL don’t add any labels. If ’n’ no labels but show the points. If equal coord add coordinates to the points. Otherwise number all points from one.

**Value**  
The ggplot object.
Note

Currently only points are checked for dominance. That is, for MILP models some nondominated points may in fact be dominated by a segment.

Author(s)

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Examples

dat <- data.frame(z1=c(12,14,16,18,18,18,14,15,15), z2=c(18,16,12,4,2,6,14,14,16))
points <- addNDSet(dat, crit = "min", keepDom = TRUE)
plotNDSet2D(points, crit = "min", addTriangles = TRUE)
plotNDSet2D(points, crit = "min", addTriangles = FALSE)
plotNDSet2D(points, crit = "min", addTriangles = TRUE, addHull = FALSE)
points <- addNDSet(dat, crit = "max", keepDom = TRUE)
plotNDSet2D(points, crit = "max", addTriangles = TRUE)
plotNDSet2D(points, crit = "max", addHull = FALSE)

plotPlane3D

Plot a plane in 3D.

Description

Plot a plane in 3D.

Usage

plotPlane3D(
  normal,
  point = NULL,
  offset = 0,
  useShade = TRUE,
  useLines = FALSE,
  usePoints = FALSE,
  ...
)

Arguments

normal Normal to the plane.
point A point on the plane.
offset The offset of the plane (only used if point = NULL).
useShade Plot shade of the plane.
useLines Plot lines inside the plane.
usePoints Plot point shapes inside the plane.
Further arguments passed on the the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:

- **argsPlanes3d**: A list of arguments for `rgl::planes3d()` used when `useShade = TRUE`.
- **argsLines**: A list of arguments for `rgl::persp3d()` when `useLines = TRUE`. Moreover, the list may contain `lines`: number of lines.

### Value

NULL (invisible)

### Examples

```r
ini3D(argsPlot3d = list(xlim = c(-1,10), ylim = c(-1,10), zlim = c(-1,10)) )
plotPlane3D(c(1,1,1), point = c(1,1,1))
plotPoints3D(c(1,1,1))
plotPlane3D(c(1,2,1), point = c(2,2,2), argsPlanes3d = list(color="red"))
plotPoints3D(c(2,2,2))
plotPlane3D(c(2,1,1), offset = -6, argsPlanes3d = list(color="blue"))
plotPlane3D(c(2,1,1), argsPlanes3d = list(color="green"))
finalize3D()
ini3D(argsPlot3d = list(xlim = c(-1,10), ylim = c(-1,10), zlim = c(-1,10)) )
plotPlane3D(c(1,1,1), point = c(1,1,1), useLines = TRUE, useShade = TRUE)
ids <- plotPlane3D(c(1,2,1), point = c(2,2,2), argsLines = list(col="blue", lines = 100), useLines = TRUE)
finalize3D()
# pop3d(id = ids) # remove last plane
```

---

**plotPoints3D**  
*Plot points in 3D.*

### Description

Plot points in 3D.

### Usage

```r
plotPoints3D(pts, addText = FALSE, ...)
```

### Arguments

- **pts**: A vector or matrix with the points.
- **addText**: Add text to the points. Currently `coord` (coordinates), `rownames` (rownames) and both supported or a vector with the text.
Further arguments passed on the the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:

- `argsPlot3d`: A list of arguments for `rgl::plot3d`.
- `argsPch3d`: A list of arguments for `rgl::pch3d`.
- `argsText3d`: A list of arguments for `rgl::text3d`.

**Value**

Object ids (invisible).

**Examples**

```r
ini3D()
pts<-matrix(c(1,1,1,5,5,5), ncol = 3, byrow = TRUE)
plotPoints3D(pts)
plotPoints3D(c(2,3,3), argsPlot3d = list(col = "red", size = 10))
plotPoints3D(c(3,2,3), argsPlot3d = list(col = "blue", size = 10, type="p"))
plotPoints3D(c(1.5,1.5,1.5), argsPlot3d = list(col = "blue", size = 10, type="p"))
plotPoints3D(c(2,2,2, 1,1,1), addText = "coord")
ids <- plotPoints3D(c(3,3,3, 4,4,4), addText = "rownames")
finalize3D()
rgl::rglwidget()
# pop3d(ids) # remove the last again
```

---

**plotPolygon3D**

*Plot a polygon.*

**Description**

Plot a polygon.

**Usage**

```r
plotPolygon3D(
    pts,
    useShade = TRUE,
    useLines = FALSE,
    usePoints = FALSE,
    useFrame = TRUE,
    ...
)
```
Arguments

- **pts**  
  Vertices.
- **useShade**  
  Plot shade of the polygon.
- **useLines**  
  Plot lines inside the polygon.
- **usePoints**  
  Plot point shapes inside the polygon.
- **useFrame**  
  Plot a frame around the polygon.

... Further arguments passed on the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:

- **argsShade**  
  A list of arguments for `rgl::polygon3d` (n > 4 vertices), `rgl::triangles3d()` (n = 3 vertices) and `rgl::quads3d()` (n = 4 vertices) if `useShade = TRUE`.
- **argsFrame**  
  A list of arguments for `rgl::lines3d` if `useFrame = TRUE`.
- **argsPoints**  
  A list of arguments for `rgl::shade3d` if `usePoints = TRUE`. It is important to give a texture using `texture`. A texture can be set using `getTexture()`.
- **argsLines**  
  A list of arguments for `rgl::persp3d()` when `useLines = TRUE`. Moreover, the list may contain `lines`: number of lines.

Value

Object ids (invisible).

Examples

```r
pts <- data.frame(x = c(1,0,0,0.4), y = c(0,1,0,0.3), z = c(0,0,1,0.3))
pts <- data.frame(x = c(1,0,0), y = c(0,1,0), z = c(0,0,1))

ini3D()
plotPolygon3D(pts)
finalize3D()

ini3D()
plotPolygon3D(pts, argsShade = list(color = "red", alpha = 1))
finalize3D()

ini3D()
plotPolygon3D(pts, useFrame = TRUE, argsShade = list(color = "red", alpha = 0.5),
              argsFrame = list(color = "green"))
finalize3D()

ini3D()
plotPolygon3D(pts, useFrame = TRUE, useLines = TRUE, useShade = TRUE,
              argsShade = list(color = "red", alpha = 0.2),
              argsLines = list(color = "blue"))
finalize3D()

ini3D()
```
```
ids <- plotPolygon3D(pts, usePoints = TRUE, useFrame = TRUE,
    argsPoints = list(texture = getTexture(pch = 16, cex = 20)))
finalize3D()
# pop3d(id = ids) # remove object again

# In general you have to finetune size and numbers when you use textures
# Different pch
for (i in 0:3) {
    fname <- getTexture(pch = 15+i, cex = 30)
    ini3D(TRUE)
    plotPolygon3D(pts, usePoints = TRUE, argsPoints = list(texture = fname))
    finalize3D()
}

# Size of pch
for (i in 1:4) {
    fname <- getTexture(pch = 15+i, cex = 10 * i)
    ini3D(TRUE)
    plotPolygon3D(pts, usePoints = TRUE, argsPoints = list(texture = fname))
    finalize3D()
}

# Number of pch
fname <- getTexture(pch = 16, cex = 20)
for (i in 1:4) {
    ini3D(TRUE)
    plotPolygon3D(pts, usePoints = TRUE,
        argsPoints = list(texture = fname, texcoords = rbind(pts$x, pts$y, pts$z)*5*i))
    finalize3D()
}
```
plotPolytope

crit = "max",
faces = type,
plotFaces = TRUE,
plotFeasible = TRUE,
plotOptimum = FALSE,
latex = FALSE,
labels = NULL,
...
)

Arguments

A        The constraint matrix.
b        Right hand side.
obj      A vector with objective coefficients.
type     A character vector of same length as number of variables. If entry k is 'i' variable
          k must be integer and if 'c' continuous.
nonneg   A boolean vector of same length as number of variables. If entry k is TRUE then
          variable k must be non-negative.
crit     Either max or min (only used if add the iso-profit line)
faces    A character vector of same length as number of variables. If entry k is 'i' variable
          k must be integer and if 'c' continuous. Useful if e.g. want to show the linear
          relaxation of an IP.
plotFaces If True then plot the faces.
plotFeasible If True then plot the feasible points/segments (relevant for IPLP/MILP).
plotOptimum Show the optimum corner solution point (if alternative solutions only one is
          shown) and add the iso-profit line.
llatex    If True make latex math labels for TikZ.
labels If NULL don’t add any labels. If ’n’ no labels but show the points. If equal coord
          add coordinates to the points. Otherwise number all points from one.

... If 2D, further arguments passed on the the ggplot plotting functions. This must be
done as lists. Currently the following arguments are supported:
  • argsFaces: A list of arguments for plotHull2D.
  • argsFeasible: A list of arguments for ggplot2 functions:
    – geom_point: A list of arguments for ggplot2::geom_point.
    – geom_line: A list of arguments for ggplot2::geom_line.
  • argsLabels: A list of arguments for ggplot2 functions:
    – geom_text: A list of arguments for ggplot2::geom_text.
  • argsOptimum:
    – geom_point: A list of arguments for ggplot2::geom_point.
    – geom_abline: A list of arguments for ggplot2::geom_abline.
    – geom_label: A list of arguments for ggplot2::geom_label.
  • argsTheme: A list of arguments for ggplot2::theme.
If 3D further arguments passed on the the RGL plotting functions. This must be done as lists. Currently the following arguments are supported:

- `argsAxes3d`: A list of arguments for `rgl::axes3d`.
- `argsPlot3d`: A list of arguments for `rgl::plot3d` to open the RGL window.
- `argsTitle3d`: A list of arguments for `rgl::title3d`.
- `argsFaces`: A list of arguments for `plotHull3D`.
- `argsFeasible`: A list of arguments for RGL functions:
  - `points3d`: A list of arguments for `rgl::points3d`.
  - `segments3d`: A list of arguments for `rgl::segments3d`.
  - `triangles3d`: A list of arguments for `rgl::triangles3d`.
- `argsLabels`: A list of arguments for RGL functions:
  - `points3d`: A list of arguments for `rgl::points3d`.
  - `text3d`: A list of arguments for `rgl::text3d`.
- `argsOptimum`: A list of arguments for RGL functions:
  - `points3d`: A list of arguments for `rgl::points3d`.

### Value

If 2D a `ggplot` object. If 3D a RGL window with the 3D plot.

### Note

The feasible region defined by the constraints must be bounded (i.e. no extreme rays) otherwise you may see strange results.

### Author(s)

Lars Relund <lars@relund.dk>

### Examples

#### 2D examples ####

# Define the model max/min coeff*x st. Ax<=b, x>=0
A <- matrix(c(-3,2,2,4,9,10), ncol = 2, byrow = TRUE)
b <- c(3,27,90)
obj <- c(7.75, 10)
## LP model
# The polytope with the corner points
plotPolytope(A,
b,
obj,
type = rep("c", ncol(A)),
crit = "max",
faces = rep("c", ncol(A)),
plotFaces = TRUE,
plotFeasible = TRUE,
```
plotPolytope(
  plotOptimum = FALSE,
  labels = NULL,
  argsFaces = list(argsGeom_polygon = list(fill = "red"))
)
# With optimum and labels:
plotPolytope(
  A,
  b,
  obj,
  type = rep("c", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "coord",
  argsOptimum = list(lty="solid")
)
# Minimize:
plotPolytope(
  A,
  b,
  obj,
  type = rep("c", ncol(A)),
  crit = "min",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "n"
)
# Note return a ggplot so can e.g. add other labels on e.g. the axes:
p <- plotPolytope(
  A,
  b,
  obj,
  type = rep("c", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "coord"
)
p + ggplot2::xlab("x") + ggplot2::ylab("y")

# More examples

## LP-model with no non-negativity constraints
A <- matrix(c(-3, 2, 2, 4, 9, 10, 1, -2), ncol = 2, byrow = TRUE)
b <- c(3, 27, 90, 2)
obj <- c(7.75, 10)
plotPolytope(
```
A,
b,
obj,
type = rep("c", ncol(A)),
nonneg = rep(FALSE, ncol(A)),
crit = "max",
faces = rep("c", ncol(A)),
plotFaces = TRUE,
plotFeasible = TRUE,
plotOptimum = FALSE,
labels = NULL)

## The package don't plot feasible regions that are unbounded e.g if we drop the 2 and 3 constraint
A <- matrix(c(-3,2), ncol = 2, byrow = TRUE)
b <- c(3)
obj <- c(7.75, 10)
# Wrong plot
plotPolytope(
  A,
b,
  obj,
  type = rep("c", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = FALSE,
  labels = NULL)

# One solution is to add a bounding box and check if the bounding box is binding
A <- rbind(A, c(1,0), c(0,1))
b <- c(b, 10, 10)
plotPolytope(
  A,
b,
  obj,
  type = rep("c", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = FALSE,
  labels = NULL)

## ILP model
A <- matrix(c(-3,2,2,4,9,10), ncol = 2, byrow = TRUE)
b <- c(3,27,90)
obj <- c(7.75, 10)
# ILP model with LP faces:
plotPolytope(
  A,
  b,
  obj,
  type = rep("i", ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "coord",
  argsLabels = list(size = 4, color = "blue"),
  argsFeasible = list(color = "red", size = 3)
)

# ILP model with IP faces:
plotPolytope(
  A,
  b,
  obj,
  type = rep("i", ncol(A)),
  crit = "max",
  faces = rep("i", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "coord"
)

## MILP model
A <- matrix(c(-3, 2, 4, 9, 10), ncol = 2, byrow = TRUE)
b <- c(3, 27, 90)
obj <- c(7.75, 10)
# Second coordinate integer
plotPolytope(
  A,
  b,
  obj,
  type = c("c", "i"),
  crit = "max",
  faces = c("c", "i"),
  plotFaces = FALSE,
  plotFeasible = TRUE,
  plotOptimum = TRUE,
  labels = "coord",
  argsFeasible = list(color = "red")
)

# First coordinate integer and with LP faces:
plotPolytope(
  A,
  b,
  obj,
plotPolytope

type = c("i", "c"),
crit = "max",
faces = c("c", "c"),
plotFaces = TRUE,
plotFeasible = TRUE,
plotOptimum = TRUE,
labels = "coord"
)

# First coordinate integer and with LP faces:
plotPolytope(
A,
b,
obj,
type = c("i", "c"),
crit = "max",
faces = c("i", "c"),
plotFaces = TRUE,
plotFeasible = TRUE,
plotOptimum = TRUE,
labels = "coord"
)

#### 3D examples ####

# Ex 1
view <- matrix(c(-0.412063330411911, -0.228006735444069, 0.882166087627411, 0, 0.910147845745087,
 -0.0574885793030262, 0.410274744033813, 0, -0.042830865830183, 0.97196090221405,
 0.231208890676498, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix(c(3, 2, 5,
 2, 1, 1,
 1, 1, 3,
 5, 2, 4), nc = 3, byrow = TRUE)
b <- c(55, 26, 30, 57)
obj <- c(20, 10, 15)
# LP model
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord")
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord",
 argsFaces = list(drawLines = FALSE, argsPolygon3d = list(alpha = 0.95)),
 argsLabels = list(points3d = list(color = "blue"))
# ILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "i"), plotOptimum = TRUE, obj = obj)
# MILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "c", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("c", "i", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "c"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "c"), plotFaces = FALSE)
plotPolytope(A, b, type = c("i", "c", "c"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "i", "c"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "c", "i"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)

# Ex 2
view <- matrix(c(-0.812462985515594, -0.029454167932272, 0.582268416881561, 0, 0.579295456409454, 0.55, -0.153386667370796, 0.800555109977722, 0, 0.0657325685024261, 0.987727105617523, 0.14168381690979, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix(c(1, 1, 1,
             3, 0, 1), nc = 3, byrow = TRUE)
b <- c(10, 24)
obj <- c(20, 10, 15)
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord")
# ILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "i"), plotOptimum = TRUE, obj = obj)
# MILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "c", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "c"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "c"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, type = c("i", "i", "c"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "c", "i"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "i", "c"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "c", "i"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)

# Ex 3
view <- matrix(c(0.976349174976349, -0.202332556247711, 0.0761845782399178, 0, 0.0903248339891434, 0.0, 0.976349174976349, -0.202332556247711, 0.0761845782399178, 0, 0.0903248339891434, 0.701892614364624, 0.706531345844269, 0, -0.196427255868912, -0.682940244674683, 0.70356869675708, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
A <- matrix(c(-1, 1, 0,
             1, 4, 0,
             2, 1, 0,
             3, -4, 0,
             0, 0, 4), nc = 3, byrow = TRUE)
b <- c(5, 45, 27, 24, 10)
obj <- c(5, 45, 15)
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord")
# ILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "i"), plotOptimum = TRUE, obj = obj)
# MILP model
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "c", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("c", "i", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "i"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, faces = c("c", "c", "c"), type = c("i", "i", "c"), plotOptimum = TRUE, obj = obj)
plotPolytope(A, b, type = c("i", "c", "i"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "i", "c"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)
plotPolytope(A, b, type = c("c", "c", "i"), plotOptimum = TRUE, obj = obj, plotFaces = FALSE)

# Ex 4
Plot the polytope (bounded convex set) of a linear mathematical program

**Description**
Plot the polytope (bounded convex set) of a linear mathematical program

**Usage**
```r
plotPolytope2D(A,
               b,
               obj = NULL,
               type = rep("c", ncol(A)),
               nonneg = rep(TRUE, ncol(A)),
               crit = "max",
               faces = rep("c", ncol(A)),
               plotFaces = TRUE,
               plotFeasible = TRUE,
               plotOptimum = TRUE)
```

```r
view <- matrix(c(-0.452365815639496, -0.446501553058624, 0.77201122045517, 0, 0.886364221572876,
                 -0.320795893669128, 0.333835482597351, 0, 0.0986008867621422, 0.835299551486969,
                 0.540881276130676, 0, 0, 0, 0, 1), nc = 4)
loadView(v = view)
Ab <- matrix(c(1, 1, 2, 5,
              2, -1, 0, 3,
              -1, 2, 1, 3,
              0, -3, 5, 2,
              # 1, 0, 1, 0, 4,
              # 1, 0, 0, 4), nc = 4, byrow = TRUE)
A <- Ab[,1:3]
b <- Ab[,4]
obj <- c(1,1,3)
plotPolytope2D()
```
plotPolytope2D

latex = FALSE,
labels = NULL,
...
)

Arguments

A       The constraint matrix.
b       Right hand side.
obj     A vector with objective coefficients.
type    A character vector of same length as number of variables. If entry k is 'i' variable k must be integer and if 'c' continuous.
nonneg  A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.
crit    Either max or min (only used if add the iso-profit line)
faces   A character vector of same length as number of variables. If entry k is 'i' variable k must be integer and if 'c' continuous. Useful if e.g. want to show the linear relaxation of an IP.
plotFaces If True then plot the faces.
plotFeasible If True then plot the feasible points/segments (relevant for ILP/MILP).
plotOptimum Show the optimum corner solution point (if alternative solutions only one is shown) and add the iso-profit line.
latex   If True make latex math labels for TikZ.
labels  If NULL don’t add any labels. If ‘n’ no labels but show the points. If equal coord add coordinates to the points. Otherwise number all points from one.
...
Further arguments passed on the the ggplot plotting functions. This must be done as lists. Currently the following arguments are supported:
  • argsFaces: A list of arguments for plotHull2D.
  • argsFeasible: A list of arguments for ggplot functions:
    – geom_point: A list of arguments for ggplot2::geom_point.
    – geom_line: A list of arguments for ggplot2::geom_line.
  • argsLabels: A list of arguments for ggplot functions:
    – geom_text: A list of arguments for ggplot2::geom_text.
  • argsOptimum:
    – geom_point: A list of arguments for ggplot2::geom_point.
    – geom_abline: A list of arguments for ggplot2::geom_abline.
    – geom_label: A list of arguments for ggplot2::geom_label.
  • argsTheme: A list of arguments for ggplot2::theme.

Value

A ggplot object.
Note
In general use `plotPolytope()` instead of this function. The feasible region defined by the constraints must be bounded otherwise you may see strange results.

Author(s)
Lars Relund <lars@relund.dk>

See Also
`plotPolytope()` for examples.

---

### plotPolytope3D

*Plot the polytope (bounded convex set) of a linear mathematical program*

#### Description
Plot the polytope (bounded convex set) of a linear mathematical program

#### Usage
```r
plotPolytope3D(
  A,            # The constraint matrix.
  b,            # Right hand side.
  obj = NULL,   # A vector with objective coefficients.
  type = rep("c", ncol(A)),
  nonneg = rep(TRUE, ncol(A)),
  crit = "max",
  faces = rep("c", ncol(A)),
  plotFaces = TRUE,
  plotFeasible = TRUE,
  plotOptimum = FALSE,
  latex = FALSE,
  labels = NULL,
  ...            # Additional arguments
)
```

#### Arguments
- `A`: The constraint matrix.
- `b`: Right hand side.
- `obj`: A vector with objective coefficients.
- `type`: A character vector of same length as number of variables. If entry k is ’i’ variable k must be integer and if ’c’ continuous.
nonneg
A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.
crit
Either max or min (only used if add the iso-profit line)
faces
A character vector of same length as number of variables. If entry k is 'i' variable k must be integer and if 'c' continuous. Useful if e.g. want to show the linear relaxation of an IP.
plotFaces
If True then plot the faces.
plotFeasible
If True then plot the feasible points/segments (relevant for ILP/MILP).
plotOptimum
Show the optimum corner solution point (if alternative solutions only one is shown) and add the iso-profit line.
latex
If NULL don't add any labels. If 'n' no labels but show the points. If equal coord add coordinates to the points. Otherwise number all points from one.
labels
If NULL don't add any labels. If 'n' no labels but show the points. If equal coord add coordinates to the points. Otherwise number all points from one.

Value
A RGL window with 3D plot.

Note
In general use `plotPolytope()` instead of this function. The feasible region defined by the constraints must be bounded otherwise you may see strange results.

Author(s)
Lars Relund <lars@relund.dk>

See Also
`plotPolytope()` for examples.
plotRectangle3D

Plot a rectangle defined by two corner points.

Description

The rectangle is defined by \( \{ x \mid a \leq x \leq b \} \) where \( a \) is the minimum values and \( b \) is the maximum values.

Usage

plotRectangle3D(a, b, ...)

Arguments

- **a**: A vector of length 3.
- **b**: A vector of length 3.
- **...**: Further arguments passed on the the RGL plotting functions. This must be done as lists (see examples). Currently the following arguments are supported:
  - **argsPlot3d**: A list of arguments for \texttt{rgl::plot3d}.
  - **argsSegments3d**: A list of arguments for \texttt{rgl::segments3d}.
  - **argsPolygon3d**: A list of arguments for \texttt{rgl::polygon3d}.
  - **argsShade3d**: A list of arguments for \texttt{rgl::shade3d}.

Value

Object ids (invisible).

Examples

```r
ini3D()
plotRectangle3D(c(0,0,0), c(1,1,1))
plotRectangle3D(c(1,1,1), c(4,4,3), drawPoints = TRUE, drawLines = FALSE,
               argsPlot3d = list(size=2, type="s", alpha=0.3))
ids <- plotRectangle3D(c(2,2,2), c(3,3,2.5), argsPolygon3d = list(alpha = 1) )
finalize3D()
# pop3d(id = ids) remove last object
```
plotTeX3D

Plot TeX at a position.

Description

Plot TeX at a position.

Usage

plotTeX3D(
  x,
  y,
  z,
  tex,
  cex = graphics::par("cex"),
  fixedSize = FALSE,
  size = 480,
  ...
)

Arguments

x  Coordinate.
y  Coordinate.
z  Coordinate.
tex  TeX string.
cex  Expansion factor (you properly have to fine tune it).
fixedSize  Fix the size of the object (no scaling when zoom).
size  Size of the generated png.
...  Arguments passed on to rgl::sprites3d() and texToPng().

Value

The shape ID of the displayed object is returned.

Examples

## Not run:
tex0 <- "$\mathbb{R}_{\geqq}\$"
tex1 <- "\LaTeX"
tex2 <- "This is a title"
ini3D(argsPlot3d = list(xlim = c(0, 2), ylim = c(0, 2), zlim = c(0, 2)))
plotTeX3D(0.75,0.75,0.75, tex0)
plotTeX3D(0.5,0.5,0.5, tex0, cex = 2)
plotTeX3D(1,1,1, tex2)
finalize3D()
**plotTitleTeX3D**

*Draw boxes, axes and other text outside the data using TeX strings.*

**Description**

- **Description:**
  - Draw boxes, axes and other text outside the data using TeX strings.

**Usage**

```r
plotTitleTeX3D(
  main = NULL,
  sub = NULL,
  xlab = NULL,
  ylab = NULL,
  zlab = NULL,
  line = NA,
  ...
)
```

**Arguments**

- **main**: The main title for the plot.
- **sub**: The subtitle for the plot.
- **xlab**: The axis labels for the plot.
- **ylab**: The axis labels for the plot.
- **zlab**: The axis labels for the plot.
- **line**: The “line” of the plot margin to draw the label on.
- **...**: Additional parameters which are passed to `plotMTeX3D`.

**Details**

The rectangular prism holding the 3D plot has 12 edges. They are identified using 3 character strings. The first character (x', y', or z') selects the direction of the axis. The next two characters are each - or +, selecting the lower or upper end of one of the other coordinates. If only one or two characters are given, the remaining characters default to -'. For example, `edge = 'x+'` draws an x-axis at the high level of y and the low level of z.

By default, axes3d uses the `bbox3d` function to draw the axes. The labels will move so that they do not obscure the data. Alternatively, a vector of arguments as described above may be used, in which case fixed axes are drawn using `axis3d`.

If `pos` is a numeric vector of length 3, `edge` determines the direction of the axis and the tick marks, and the values of the other two coordinates in `pos` determine the position. See the examples.
The object IDs of objects added to the scene.

Examples

```r
## Not run:
in3D(argsPlot3d = list(xlim = c(0, 2), ylim = c(0, 2), zlim = c(0, 2)))
plotTitleTeX3D(main = "\LaTeX", sub = "subtitle $\alpha$",
              xlab = "$x^1_2$", ylab = "$\beta$", zlab = "$x\cdot y$")
finalize3D()
## End(Not run)
```

Description

To size of the png file.

Usage

```r
pngSize(png)
```

Arguments

- `png` Png file name.

Value

A list with width and height.

saveView

Help function to save the view angle for the RGL 3D plot

Description

Help function to save the view angle for the RGL 3D plot

Usage

```r
saveView(fname = "view.RData", overwrite = FALSE, print = FALSE)
```
Arguments

fname The file name of the view.
overwrite Overwrite existing file.
print Print the view so can be copied to R code (no file is saved).

Value

NULL (invisible).

Note

Only save if the file name don’t exists.

Author(s)

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Examples

view <- matrix( c(-0.412063330411911, -0.228006735444069, 0.882166087627411, 0,
0.910147845745087, -0.0574885793030262, 0.410274744033813, 0, -0.042830865830183,
0.97196090221405, 0.231208890676498, 0, 0, 0, 0, 1), nc = 4)

loadView(v = view)
A <- matrix( c(3, 2, 5, 2, 1, 1, 1, 1, 3, 5, 2, 4), nc = 3, byrow = TRUE)
b <- c(55, 26, 30, 57)
obj <- c(20, 10, 15)
plotPolytope(A, b, plotOptimum = TRUE, obj = obj, labels = "coord")

# Try to modify the angle in the RGL window
saveView(print = TRUE) # get the view angle to insert into R code

slices Find all corner points in the slices define for each fixed integer combination.

Description

Find all corner points in the slices define for each fixed integer combination.
Usage

slices(
    A,
    b,
    type = rep("c", ncol(A)),
    nonneg = rep(TRUE, ncol(A)),
    collapse = FALSE
)

Arguments

A            The constraint matrix.
b            Right hand side.
type         A character vector of same length as number of variables. If entry k is 'i' variable k must be integer and if 'c' continuous.
nonneg       A boolean vector of same length as number of variables. If entry k is TRUE then variable k must be non-negative.
collapse     Collapse list to a data frame with unique points.

Value

A list with the corner points (one entry for each slice).

Examples

A <- matrix( c(3, -2, 1, 2, 4, -2, -3, 2, 1), nc = 3, byrow = TRUE)
b <- c(10,12,3)
slices(A, b, type=c("i","c","i"))

A <- matrix(c(9,10,2,4,-3,2), ncol = 2, byrow = TRUE)
b <- c(90,27,3)
slices(A, b, type=c("c","i"), collapse = TRUE)

texToPng

Convert LaTeX to a png file

Description

Convert LaTeX to a png file

Usage

texToPng(
    tex,
    width = NULL,
    height = NULL,
Arguments

- **tex**: TeX string. Remember to escape backslash with \\.
- **width**: Width of the png.
- **height**: Height of the png (width are ignored).
- **dpi**: Dpi of the png. Not used if width or height are specified.
- **viewPng**: View the result in the plots window.
- **fontsize**: Front size used in the LaTeX document.
- **calcM**: Estimate 1 em in pixels in the resulting png.
- **crop**: Call command line program `pdfcrop` (must be installed).

Value

The filename of the png or a list if calcM = TRUE.

Examples

```r
## Not run:
tex <- "$\mathbb{R}_{\geqq}\$"
texToPng(tex, viewPng = TRUE)
texToPng(tex, fontsize = 20, viewPng = TRUE)
texToPng(tex, height = 50, fontsize = 10, viewPng = TRUE)
texToPng(tex, height = 50, fontsize = 50, viewPng = TRUE)
tex <- "MMM"
texToPng(tex, dpi=72, calcM = TRUE)
texToPng(tex, width = 100, calcM = TRUE)
f <- texToPng(tex, dpi=300)
pngSize(f)

## End(Not run)
```
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