

# Package ‘knotR’

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**Type** Package

**Title** Knot Diagrams using Bezier Curves

**Version** 1.0-4

**Depends** R (>= 2.10)

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**LazyData** TRUE

**Description** Makes visually pleasing diagrams of knot projections using optimized Bezier curves.

**License** GPL-2

**NeedsCompilation** no

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knotR-package	<i>Knot Diagrams using Bezier Curves</i>
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**Description**

Makes visually pleasing diagrams of knot projections using optimized Bezier curves.

**Details**

The DESCRIPTION file:

```
Package:      knotR
Type:         Package
Title:        Knot Diagrams using Bezier Curves
Version:      1.0-4
Authors@R:    person(given=c("Robin", "K. S."), family="Hankin", role = c("aut","cre"), email="hankin.robin@gmail.com",
Depends:      R (>= 2.10)
Maintainer:   Robin K. S. Hankin <hankin.robin@gmail.com>
LazyData:    TRUE
Description:  Makes visually pleasing diagrams of knot projections using optimized Bezier curves.
License:      GPL-2
Author:       Robin K. S. Hankin [aut, cre] (<https://orcid.org/0000-0001-5982-0415>)
```

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knots	Optimized knots
overunder	Functionality for specifying overstrands and understrands
reader	Reading and writing svg files
symmetrize	Symmetry and knots
utilities	Various utilities for knots

The package contains a large number of knots, optimized for visual appearance in the sense that the

knot path is nice and smooth, and strands cross at close to right angles. These can be displayed by typing

```
knotplot(k9_23)
```

at the R prompt. The package includes all prime knots up to and including 9 crossings, and a number of other interesting and attractive knots.

The package facilitates the creation and optimization of new knots. The basic workflow is to create an .svg file in inkscape comprising a single closed path (that is, the first and last node are the same point). Control nodes should all be symmetrical. Many examples of correctly formatted .svg files are given in the `inst/` directory.

The best way to reproduce a knot from an image of its projection is to fire up inkscape, then *import* the image into inkscape, resize and rotate as desired, then follow the string with the ‘Bezier curves and straight lines’ tool (also called the ‘pen tool’ by Kirsanov; the keyboard shortcut is shift-F6). Use 1-2 nodes per segment, or 3 nodes for longer or more visually prominent segments.

Keep the lines straight at first. Close the path by making a final click on the initial node; now you have a closed polygon, which will self-intersect at the path crossing points. To smoothen the path, select the ‘edit paths by node’ tool (shift-F2), then convert the corner nodes of the path to symmetric Bezier nodes (‘make selected nodes symmetric’). You can then tweak the path by moving the control nodes about with the mouse. Be aware that adding or deleting nodes changes the adjacent nodes to asymmetrical Bezier control points; make them symmetric by selecting all nodes (‘Ctrl-A’), then hit the ‘make selected nodes symmetric’ button. Do this frequently to avoid confusion.

An .svg file may be imported into R using the `reader()` function, which creates an `inkscape` object. This represents the *path* of the knot: it does not include over and under information. The package assumes that inkscape uses absolute coordinates (as opposed to relative coordinates); see `reader.Rd` for more information.

The package provides four classes of objects that specify the path of a knot: `inkscape`, `minobj`, `controlpoints`, and `knotvec`. These four classes have different uses, and objects may be converted from one form to another by using functions such as `as.minobj()`, documented at `as.Rd` and `utilities.Rd`.

A knot requires information on which strands pass over or under which other strands; full documentation at `?overunder`.

Knots sometimes have symmetry constraints such as horizontal or vertical symmetry, or rotational symmetry. Symmetry is imposed by using the `symmetrize()` function: this takes a knot path (coerced to `minobj` form) and a symmetry object. Symmetry objects are created with function `symmetry_object()`, which takes a knot path and a series of matrices and vectors that specify the symmetry of the knot.

### Author(s)

NA

Maintainer: Robin K. S. Hankin <hankin.robin@gmail.com>

### Examples

```
a <- reader(system.file("7_6.svg",package="knotR"))
knotplot2(a) # shows curvature

# Now use text=TRUE to display strand numbers so you can figure out the
# overunder relations:

knotplot2(a,text=TRUE,lwd=1)

ou76 <- matrix(c(
  12,01,
  02,11,
  07,03,
  04,15,
  16,06,
  14,08,
  10,13
),byrow=TRUE,ncol=2)

# Now we can do a proper knot plot:

knotplot(a,ou76)

# To symmetrize a knot we use the symmetry functionality of the knot:

a <- reader(system.file("3_1_not_symmetric.svg",package="knotR"))

knotplot2(a,seg=TRUE,text=TRUE,lwd=1,node=TRUE)

# First specify the vertical symmetry:

Mver <- matrix(c(
  08,10,
  07,11,
  02,04,
  01,05,
  12,06
),ncol=2,byrow=TRUE)

# Then the rotational symmetry:

Mrot <- matrix(c(
  09,05,01,
  10,06,02,
  08,04,12
),byrow=TRUE,ncol=3)

# Now the overunder information:

ou31 <- matrix(c(
  03,08,
  11,04,
  07,12
),byrow=TRUE,ncol=2)
```

```

# create a symmetry object:

sym31 <- symmetry_object(a, Mver=Mver,xver=c(9,3),Mrot=Mrot)

knotplot(symmetrize(a,sym31),ou31)

# Symmetric-- but ugly as a burglar's bulldog.

# to beautify, either use the knotoptim() function, or do it by hand:

objective <- function(m) {badness(make_minobj_from_minsymvec(m, sym31))}
startval <- make_minsymvec_from_minobj(as.minobj(a),sym31)

## Not run:

# Following examples take a long time to run.
# nlm() is the best optimization method, I think. Limit to 1 iteration:
o <- nlm(f=objective, p=startval, iterlim=1)

# extract the evaluate:
oo <- make_minobj_from_minsymvec(o$estimate, sym31)

# create a knot:
k31_marginally_better <-
knot(x = oo, overunderobj = ou31, symobj = sym31)

# then plot it:
knotplot(k31_marginally_better)

## End(Not run)

```

---

as

*Conversions between various forms of a knot*


---

## Description

Conversions between various forms of a knot.

## Usage

```

as.knotvec(x)
as.minobj(x)
as.inkscape(x)
as.controlpoints(x)
as.minsymvec(x,symobj)

```

**Arguments**

x                    Object to be converted  
 symobj              A symmetry object

**Details**

The `as.foo()` functions are meant to be user-friendly; they use low-level functions like `make_knotvec_from_minobj()` (all of which are documented at `utilities.Rd`), which are a bit messy.

**Author(s)**

Robin K. S. Hankin

**See Also**

[utilities](#)

**Examples**

```
as.minobj(k6_2)

x <- reader(system.file("6_3.svg",package="knotR")) # x is class inkscape

as.minsymvec(x,symmetry_object(k6_3)) # as.minsymvec() needs a symmetry object

as.controlpoints(x)

as.knotvec(x)
```

---

badness

*Badness of knots*

---

**Description**

Various functions that calculate different aspects of the badness of a knot, generally with low values representing pleasing visual representations

**Usage**

```
badness(b, cpb, weights, prob=0, give=FALSE)
curvature_switching_badness(b)
curvature_consecutive_segment_switching_badness(b, ...)
midpoint_badness(b, cpb)
node_crossing_badness(b, cpb)
total_string_length(b)
total_bending_energy(b, power=2)
```

```
total_crossing_potential_energy(b, cpb)
total_crossing_angle_badness(b, cpb)
metrics(b, cpb)
always_left_badness(b)
non_crossing_strand_close_approach_badness(b, cpb)
```

### Arguments

<code>b</code>	A description of a knot, coerced to a controlpoints object
<code>cpb</code>	Optional argument containing information on crossing points (it is short for ‘crossing_points(b)’). It is time-consuming to calculate, so providing a pre-calculated value makes the code run faster
<code>prob</code>	In function <code>badness()</code> , the probability of plotting a knotplot. I used nonzero values in the early stages of developing the package: when optimizing a knot it was useful to keep tabs on the process
<code>give</code>	In function <code>badness()</code> , Boolean with default FALSE meaning to return the sum of the badnesses, and TRUE meaning to return them separately
<code>power</code>	Function <code>total_bending_energy()</code> returns the arc integral of $R^{-p}$ ; defaults to 2
<code>weights</code>	A vector of weights specifying the relative importance of the various badness measures. See details
<code>...</code>	In function <code>curvature_consecutive_segment_switching_badness()</code> , extra arguments passed to <code>integrate()</code>

### Details

Various functions that calculate different aspects of the badness of a knot, generally with low values representing pleasing visual representations. Function `badness()` returns a weighted sum of nine individual badnesses.

The list below details the values returned by `metrics()`; the description of each item is the name of corresponding weight assigned by the `weights` argument of `badness()`.

- pot** Function `total_crossing_potential_energy()` gives the potential energy of the nodes, under an inverse square force law
- ang** Function `total_crossing_angle_badness()` returns a high value if strands cross at angles far from 90 degrees. It returns the sum, over all crossings, of `bezier_angle()`
- ben** Function `total_bending_energy()` gives the total bending energy, effectively the arc integral of the reciprocal of the square of the radius of curvature
- len** Function `total_string_length()` returns  $\ell$ , the total string length. The badness is proportional to  $(\ell - 5000)^2$ . A length of 5000 corresponds to knots that look about right on a sheet of A4 paper
- mid** Function `midpoint_badness()` penalizes knots with crossing points far from the midpoint of segments
- clo** Function `node_crossing_badness()` penalizes knots with nodes too close together (compare function `total_crossing_potential_energy()`)

- swi** Function `curvature_switching_badness()` provides a penalty for consecutive segments with curvatures that switch sign. The magnitude of the penalty is zero if both curvatures are of the same sign, otherwise proportional to the square of the minimum of the maximum value of the absolute value of the positive and negative curvatures. The source code is easier to look at, honest
- con** Function `curvature_consecutive_segment_switching_badness()` penalizes knots with consecutive segments that switch curvature from positive to negative
- ncn** Function `always_left_badness()` penalizes knots that are *supposed* to curve to the left all the time (eg knot  $8_{18}$ ). The penalty is proportional to the greatest rightward curvature over the whole knot

The weights argument is nominally a vector of length 9 which is used to assign weights to different aspects of the badness of a knot.

### Value

Returns a scalar badness

### Author(s)

Robin K. S. Hankin

### See Also

[crossing](#)

### Examples

```
# use the k_infinity knot for speed:
system.time(badness(k_infinity))

cc <- crossing_points(k_infinity)
system.time(badness(k_infinity,cc))

metrics(k_infinity,cc)

## default:
badness(k_infinity, weights=c(1,1,1,1,1,1,1,1,1))

## downweight the importance of strands crossing at 90 degrees:
badness(k_infinity, weights=c(1,0.1,1,1,1,1,1,1,1))
```



**Description**

Various functionality for Bezier curves including derivatives and radius of curvature.

**Usage**

```
bezier(P, tee, n=100)
bezier_deriv(P, tee, n=100)
bezier_deriv2(P, tee, n=100)
bezier_radius(P, tee, n=100)
bezier_curvature(P, tee, n=100)
myseg(P, ...)
```

**Arguments**

P	Control points in the form of a 4 by 2 matrix with rows corresponding to $P_0$ to $P_3$
tee	Parametric variable $t$
n	Integer specifying number of points between 0 and 1 to use. Default value of 100 looks OK
...	Further arguments passed by myseg() to points()

**Details**

- Function `bezier()` returns a two column matrix with rows corresponding to the positions of the specified Bezier curve.
- Functions `bezier_deriv()` and `bezier_deriv2()` give the first and second derivatives respectively.
- Function `bezier_radius()` gives the radius of curvature.
- Functions `bezier_length()` and `bezier_bending_energy()` use numerical quadrature to give the arc length and bending energy ( $\int R^{-1} ds$ ).

**Author(s)**

Robin K. S. Hankin

**See Also**

[bezier\\_angle](#)

**Examples**

```

P <- matrix(c(0, 1, 2, 2, 2, 0, 3, 2),4,2)
xy <- bezier(P,n=100)
dx <- bezier_deriv(P,n=100)

plot(xy,asp=1)
myseg(P)

plot(xy,asp=1,cex=sqrt(rowSums(dx^2))/3.2)

plot(xy,asp=1)
segments(xy[,1],xy[,2],(xy+dx/200)[,1],(xy+dx/200)[,2])

plot(xy, asp=1,cex=bezier_radius(P,n=100)/2)

lapply(as.controlpoints(k8_9),bezier_radius)
lapply(as.controlpoints(k8_9),bezier_arclength)

```

---

bezier\_angle

*Intersection of two Bezier curves*


---

**Description**

Description of the intersection of two Bezier curves including position and angle of the point of intersection.

**Usage**

```

bezier_angle(P1, P2)
bezier_intersect(P1,P2, type='pos', ...)

```

**Arguments**

P1,P2	Control points for two Bezier curves as per bezier()
type	In function bezier_intersect(), string argument governing what exactly is to be returned; see details.
...	In function bezier_intersect(), further arguments passed to constOptim()

**Details**

Function bezier\_intersect() uses constOptim() to find the point of closest approach.

Function bezier\_angle() returns the square of the cosine of the intersection angle (so strands crossing at right angles return zero). If the strands do not intersect, then return 1. This is needed

because sometimes, strands which intersect are perturbed by the optimization routine so that they are disjoint.

In function `bezier_intersect()`, argument `type` may take the following values:

**pos** Position of intersection point

**cons** Boolean, indicating whether the strands abut; the 'intersection' point is the end of one curve and the beginning of the other

**bool** Boolean, indicating whether or not the strands actually intersect

**para** Bezier parameter  $t$  for the intersection point; actually return two parameters, one for each curve

**opt** Details of the optimization output

**all** Everything

### Note

If the curves intersect in more than one point, the behaviour of these routines is not defined.

### Author(s)

Robin K. S. Hankin

### See Also

[bezier](#)

### Examples

```
P1 <- matrix(c(1, 3, 6, 4, 7, 3, 2, 2),ncol=2)
P2 <- matrix(c(4, 5, 5, 3, 7, 2, 5, 1),ncol=2)

x1 <- bezier(P1,n=100)
x2 <- bezier(P2,n=100)

plot(x1,asp=1,xlim=c(0,8),ylim=c(0,8))
points(x2)

myseg(P1)
myseg(P2)

jj <- bezier_intersect(P1,P2)
points(x=jj[1],y=jj[2],pch=16,cex=3,col='blue')

# looks close to orthogonal, actually 82 degrees:
acos(sqrt(bezier_angle(P1,P2)))*180/pi
```

---

bezier\_find\_length     *Solve for arclength*

---

### Description

Finds the value of the Bezier parameter  $t$  that corresponds to a given arclength from the start of a Bezier curve

### Usage

```
bezier_find_length(P, len, from = 0, increasing = TRUE, give = FALSE, ...)
```

### Arguments

P	Control points in the form of a 4 by 2 matrix with rows corresponding to $P_0$ to $P_3$
from	Point from which to start measuring arc length
len	Arc length
increasing	Boolean, with default TRUE meaning to measure length towards the end, and FALSE meaning to measure in the opposite direction
give	Boolean, with TRUE meaning to return details from <code>uniroot()</code> and default FALSE meaning to give just the position on the curve
...	Further arguments passed to <code>uniroot()</code>

### Details

The function just uses `uniroot()` to find the appropriate value of  $t$ .

### Author(s)

Robin K. S. Hankin

### See Also

[bezier\\_integral](#)

### Examples

```
P <- matrix(c(1, 3, 6, 4, 7, 3, 2, 2), ncol=2)
bezier_find_length(P,5)
```

---

bezier\_integrals      *Arcwise integrals over Bezier curves*


---

**Description**

Various integrals over Bezier curves such as total arc length and bending energy

**Usage**

```
bezier_arclength(P, t1=0,t2=1,give=FALSE,...)
bezier_bending_energy(P, t1=0,t2=1, give=FALSE, power=2, ...)
```

**Arguments**

P	Control points in the form of a 4 by 2 matrix with rows corresponding to $P_0$ to $P_3$
give	Boolean, with TRUE meaning to return more information and default FALSE meaning to return just the value of the integration as estimated by the numerical routine
power	Function <code>bezier_bending_energy()</code> returns bending energy is $\int_S \frac{ds}{R^{\text{power}}}$ , where $R = R(s)$ is the radius of curvature. If <code>power = 2</code> this corresponds to the Eulerian bending energy of a flexible beam
t1, t2	In function <code>bezier_arclength()</code> , the values of t to start and end the integration at
...	Further arguments passed to <code>integrate()</code>

**Details**

These functions use numerical integration, specifically `integrate()`, between two specified points on a Bezier curve.

1. Function `bezier_bending_energy()` gives the and bending energy ( $\int R^{-1} ds$ ).
2. Function `bezier_arclength()` gives the arc length.

**Author(s)**

Robin K. S. Hankin

**See Also**

[bezier\\_angle](#)

## Examples

```
P <- matrix(c(0, 1, 2, 2, 2, 0, 3, 2),4,2)
bezier_arclength(P)
```

---

crossing

*Crossing Metrics for knots*

---

## Description

Various descriptions for the crossing points of a knot

## Usage

```
crossing_points(b, give_all = TRUE)
crossing_matrix(b)
crossing_strands(b)
```

## Arguments

**b** A list of Bezier control parameters, typically given by `getcontrolpoints()`

**give\_all** In function `crossing_points()`, Boolean, with TRUE meaning to give details of the strands that cross and default FALSE meaning to give just the coordinates of the crossing points

## Author(s)

Robin K. S. Hankin

## See Also

[as.controlpoints,bezier](#)

## Examples

```
crossing_points(k7_2,give_all=TRUE)
```

---

getstringpoints	Returns the coordinates of a knot's path
-----------------	--

---

### Description

Returns the coordinates of a knot's path

### Usage

```
getstringpoints(b, give_strand = FALSE, n = 100)
```

### Arguments

b	The knot path (coerced to controlpoints form)
give_strand	Boolean, with default FALSE meaning to return a two-column matrix with rows corresponding to coordinates of the knot path, and TRUE meaning to return a matrix with an additional column indicating the strand number
n	The number of points to use when constructing the Bezier curve

### Value

Returns either a two- or three- column matrix

### Note

Function knotplot() returns the points of the string too, but with NA for understrands.

### Author(s)

Robin K. S. Hankin

### See Also

[knotplot](#)

### Examples

```
plot(getstringpoints(k4_1),asp=1)

a <- getstringpoints(k11a179,TRUE)
plot(a,asp=1,col=rainbow(24)[a[,3]])

d <- 1200
plot(rbind(
  sweep(getstringpoints(k7_1),2,c(0,0)),
  sweep(getstringpoints(k7_2),2,c(0,d)),
  sweep(getstringpoints(k7_3),2,c(d,0)),
  sweep(getstringpoints(k7_4),2,c(d,d))
```

```
),asp=1,xlab='',ylab='')
```

---

head.inkscape      *Head and tail methods for inkscape objects*

---

### Description

Head and tail methods for inkscape objects

### Usage

```
## S3 method for class 'inkscape'
head(x, ...)
## S3 method for class 'inkscape'
tail(x, ...)
```

### Arguments

x                    Primary argument, an inkscape object  
 ...                  Further arguments, passed to head() or tail()

### Author(s)

Robin K. S. Hankin

### Examples

```
a <- reader(system.file("7_1.svg",package="knotR"))
head(a)
tail(a)

head(as.inkscape(k8_2))
```

---

knotoptim      *Optimization of knot appearance*

---

### Description

Optimization of knot appearance using user-definable objective functions

### Usage

```
knotoptim(svg, weights=1, symobj=NULL,
  Mver = NULL, xver = NULL, Mhor = NULL, xhor = NULL, Mrot = NULL,
  mcDonalds = FALSE, celtic = FALSE, ou, prob = 0, useNLM=TRUE, ...)
```



**Arguments**

svg	Name of an svg file to read
Mver, xver, Mhor, xhor, Mrot, mcDonalds, celtic	Arguments passed to <code>symmetry_object()</code> , specifying the symmetry of the knot
symobj	A symmetry object
ou	An overunder object
prob	The probability of plotting a knotplot; this is slow so don't make this too big
weights	A vector of weights, defaulting to all ones, passed to <code>badness()</code>
useNLM	Boolean, with default TRUE meaning to use <code>nlm()</code> and FALSE meaning to use <code>optim()</code>
...	Further arguments passed to <code>nlm()</code>

**Details**

Function `knotoptim()` is a generic optimization routine that starts from an svg file and minimizes the knot's `badness()`.

The `weights` argument is documented more fully at `badness.Rd`.

**Value**

Returns a knot object

**Author(s)**

Robin K. S. Hankin

**See Also**

[symmetry\\_object](#), [badness](#)

**Examples**

```
## Not run:      #takes too long
knotoptim(
  svg = system.file("4_1_first_draft.svg", package="knotR"),
  Mver = rbind(c(2,3),c(9,7),c(10,6),c(1,4),c(5,11)),
  xver = 8, # node on vertical axis
  ou = rbind( c(1,5), c(9,2), c(4,8),c(6,11)),
  prob = 0.1,
  iterlim = 100, print.level=2)

## End(Not run)
```

knotplot

*Plotting of knots***Description**

Routines to plot projections of knots with a wide range of user-settable options

**Usage**

```
knotplot(x, ou, gapwidth=1, n=100, lwd=8, add=FALSE, ...)
knotplot_old(x, ou, gap=20, n=100, lwd=8, add=FALSE, ...)
knotplot2(x, rainbow=FALSE, seg=FALSE, text=FALSE, cross=FALSE, ink=FALSE,
          node=FALSE, width=TRUE, all=FALSE, n=100, circ=1000,
          lwd=8, add=FALSE, ...)
```

**Arguments**

<code>x</code>	Description of a knot, coerced to a controlpoints object and a minobj object
<code>rainbow, seg, text, cross, ink, node, all, width, circ</code>	Variables controlling sundry knotplot2() features; see details
<code>ou</code>	An overunder object, useful if overunder information not included in argument <code>x</code>
<code>gap, gapwidth</code>	Variables controlling visual representation of strand crossings; see details
<code>n</code>	Number of points on each Bezier curve
<code>lwd</code>	Width of line to use
<code>add</code>	Boolean, with default FALSE meaning to set up a new plot, and TRUE meaning to just add points and lines to an existing plot
<code>...</code>	Further arguments, passed to plot() and points()

**Details**

Function knotplot() is useful for production-quality plotting of knots with crossings indicated by the understrand having a gap; function knotplot2() is more useful for development. Function knotplot\_old() is included for backward compatibility and is possibly more robust than knotplot().

Function knotplot() works by setting a suitable length of the understrand to NA which results in it not being plotted.

For knotplot():

- `overunderobj`; A two-column matrix indicating the sense of the crossing. Each row corresponds to a crossing; the first entry is the segment number of the overstrand, and the second is the understrand
- `gapwidth`; the width of the gap, measured in units of width of the string

For knotplot2():

- rainbow; use rainbow colouring for the segments
- seg; plot the Bezier nodes and handles. The positions of the nodes and handles are obtained from an object of class `controlpoints`.
- text; include the segment number on the segment
- cross; label the crossings
- ink; label the nodes with their inkscape numbering
- width; show the bending strain energy

The `gap` argument of `knotplot_old()` is the same as the `gapwidth` argument of `knotplot()` but `gap` is measured in the same units as the `plot()`.

### Author(s)

Robin K. S. Hankin

### Examples

```
knotplot(k5_1)
```

```
knotplot2(k6_1, text=TRUE, seg=TRUE, lwd=1)
```

---

knots

*Optimized knots*

---

### Description

A variety of knots with optimized forms

### Details

A selection of knots that have been optimized for visual appearance. The list makes no claims for completeness; the examples are intended to show the abilities of the package.

Knots with names like `k7_3` use the naming scheme of Rolfsen.

Knots with names like `k11n157` follow the nomenclature of the Hoste-Thistlethwaite table; ‘a’ means ‘alternating’ and ‘n’ means ‘nonalternating’.

Knot `k12a_614` is drawn from the “Table of Knot Invariants” by Livingstone and Cha.

Knot `amphichiral15` is the unique amphichiral knot with crossing number 15, due to Hoste, Thistlethwaite, and Weeks.

Knots `k12n_0411` and `k11a203` show that partial symmetry may be enforced.

Knot `k8_18` is an exceptional knot.

Knot `pretzel_p3_p5_p7_m3_m5` is drawn from a knot appearing in Bryant 2016. The notation specifies the sense (‘p’ for plus and ‘m’ for minus) of the twists.

Knot T20 is a “remarkable 20-crossing tangle”; see references

Knots k12a1202 and k12n838 are named following Lamm.

As of version 1.0-4, the complete list of knots is:

k10\_1, k10\_123, k10\_47, k10\_61, k12a1202, k12n838, k3\_1, k3\_1a, k4\_1, k4\_1a, k5\_1, k5\_2, k6\_1, k6\_2, k6\_3, k7\_1, k7\_2, k7\_3, k7\_4, k7\_5, k7\_6, k7\_7, k7\_7a, k8\_1, k8\_10, k8\_11, k8\_12, k8\_13, k8\_14, k8\_15, k8\_16, k8\_17, k8\_18, k8\_19, k8\_19a, k8\_19b, k8\_2, k8\_20, k8\_21, k8\_3, k8\_3\_90deg\_crossing, k8\_4, k8\_4a, k8\_5, k8\_6, k8\_7, k8\_8, k8\_9, k9\_1, k9\_10, k9\_11, k9\_12, k9\_13, k9\_14, k9\_15, k9\_16, k9\_17, k9\_18, k9\_19, k9\_2, k9\_20, k9\_21, k9\_22, k9\_23, k9\_23a, k9\_24, k9\_25, k9\_26, k9\_27, k9\_28, k9\_29, k9\_3, k9\_30, k9\_31, k9\_32, k9\_33, k9\_34, k9\_35, k9\_36, k9\_37, k9\_38, k9\_39, k9\_4, k9\_40, k9\_41, k9\_42, k9\_43, k9\_44, k9\_45, k9\_46, k9\_47, k9\_48, k9\_49, k9\_5, k9\_6, k9\_7, k9\_8, k9\_9, D16, T20, amphichiral15, celtic3, five\_loops, flower, four\_loops, hexknot, hexknot2, hexknot3, k\_infinity, k11a1, k11a179, k11a361, k11n157, k11n157\_morenodes, k11n22, k12n\_0242, k12n\_0411, longthin, ochiai, ornamental20, perko\_A, perko\_B, pretzel\_2\_3\_7, pretzel\_7\_3\_7, pretzel\_p3\_p5\_p7\_m3\_m5, reefknot, satellite, sum\_31\_41, three\_figure\_eights, trefoil\_of\_trefoils, triloop, unknot

## References

- K. A. Bryant, 2016. *Slice implies mutant-ribbon for odd, 5-stranded pretzel knots*, arXiv:1511.07009v2
- S. Eliahou and J. Fromentin 2017. “A remarkable 20-crossing tangle”. Arxiv, <https://arxiv.org/abs/1610.05560v2>

## Examples

```
knotplot(k3_1)
## maybe str(k3_1) ; plot(k3_1) ...
```

---

overunder

*Functionality for specifying overstrands and understrands*

---

## Description

Functionality for specifying overstrands and understrands

## Usage

```
overunder(x)
overunder(x) <- value
mirror(x)
is.sensible(overunderobj, x)
```

## Arguments

x                    A knot object  
value, overunderobj            A two-column integer matrix specifying the senses of the crossings in a knot

## Details

Function `overunder()` returns a two-column integer matrix with rows corresponding to crossing points. The first element of each row corresponds to the strand number of the overstrand and the second element corresponds to the understrand.

Function `is.sensible()` checks to see whether the overunder matrix is compatible with the knot path. For example, it checks to see whether each crossing has exactly one row, and that each row corresponds to a pair of strands that actually cross.

Function `mirror()` takes a knot and returns the knot with the senses of each crossing reversed; it is as though the knot is reflected in the plane of the projection.

## Author(s)

Robin K. S. Hankin

## See Also

[knot](#)

## Examples

```
overunder(k4_1)

par(mfcol=c(1,2))
knotplot(k4_1,gap=80)
knotplot(mirror(k4_1),gap=80)

is.sensible(overunder(k6_1),k6_1)
```

---

reader

*Reading and writing svg files*

---

## Description

Various utilities for reading and creating svg files for use with inkscape

## Usage

```
reader(filename)
write_svg(k, oldfile, safe=TRUE,
  regex1 = 'sodipodi:docname=',
  regex2 = ' *d *= *" *M.*C.*[zZ] *"')
```

**Arguments**

filename	Name of a file to be read by <code>reader()</code> ; usually an inkscape .svg file
safe	Boolean, with default TRUE meaning to save file “foo.svg” as “foo_smooth.svg” and FALSE meaning to overwrite foo.svg.
k, oldfile, regex1, regex2	Various arguments sent to <code>write_svg()</code> ; see the source code for details. Argument k is a knot, oldfile an .svg file for reference.

**Details**

Function `reader()` is the way to get started with a new knot. This takes a filename which is an .svg file created with inkscape. Instructions for creating a suitable inkscape file are given in `knotR-package.Rd`.

**Note**

Inkscape’s default is to use a mixture of absolute and relative coordinates. Function `reader()` assumes that the .svg file uses only absolute coordinates.

To ensure that only absolute coordinates are used, open the ‘SVG output’ menu in ‘inkscape preferences’ and uncheck the “Allow relative coordinates” option.

The format of .svg file is described in the W3C recommendation (2011) for Scalable Vector Graphics (SVG) 1.1, second edition.

Sometimes, `reader()` will fail with a valid .svg file if a node is sufficiently close to the x or y axis to require exponential notation (this typically happens with complicated rotational symmetry). If the file contains text like

```
...35.3635879230533 -1.323423734554e-15 , 10.3538368384142...
```

the second value is zero to numerical precision, but the text form of the number interferes with the operation of `reader()`. To deal with this we need to edit the file in a text editor and replace the offending number with an exact zero:

```
...35.3635879230533 0 , 10.3538368384142...
```

(I guess the ideal would be to incorporate some clever regexp technique into `reader()` but this turned out to be harder than I thought).

**Author(s)**

Robin K. S. Hankin

**See Also**

[utilities](#), [knotR-package](#)

**Examples**

```
## Not run:
a <- reader("6_3.svg")
b <- getcontrolpoints(a)
knotplot(a)
```

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

---

```
symmetrize
```

```
Symmetry and knots
```

---

## Description

Various functionality to impose different types of symmetry on knots

## Usage

```
force_nodes_mirror_images_LR(x,symobj)
force_nodes_mirror_images_UD(x,symobj)
force_nodes_exactly_horizontal(x,symobj)
force_nodes_exactly_vertical(x,symobj)
force_nodes_on_V_axis(x,xver)
force_nodes_on_H_axis(x,xhor)
force_nodes_rotational(x,symobj)
symmetrize(x,symobj)
tag_notneeded(x, Mver, xver, Mhor, xhor, Mrot,exact_h,exact_v)
make_minsymvec_from_minobj(x,symobj)
minsymvec(vec)
make_minobj_from_minsymvec(minsymvec,symobj)
symmetry_object(x, Mver=NULL, xver=NULL, Mhor=NULL, xhor=NULL,
Mrot=NULL, exact_h=NULL, exact_v=NULL,
mcdonalds=FALSE, celtic=FALSE, reefknot=FALSE,center_crossing=FALSE)
knot(x, overunderobj, symobj, Mver=NULL, xver=NULL, Mhor=NULL,
xhor=NULL, Mrot=NULL, mcdonalds=FALSE, celtic=FALSE,
reefknot=FALSE,center_crossing=FALSE)
```

## Arguments

x	Object coerced to class minobj
Mver, Mhor	Matrices specifying vertical (resp. horizontal) symmetry, with two columns. The rows specify pairs of symmetric nodes about a vertical (resp. horizontal) axis. Nodes specified by the first column should be on the left (resp. upper) side; these are fixed. Used by functions force_nodes_mirror_images_LR() and force_nodes_mirror_images_UD() which move the right (resp. lower) nodes and their associated handles to the positions required for exact vertical (resp. horizontal) symmetry
Mrot	A matrix specifying rotational symmetry. Each row corresponds to a set of nodes in a rotational relationship. The number of columns specifies the order of the rotational symmetry. The first column corresponds to nodes whose position is fixed. Used by force_nodes_rotational(), which also moves handles appropriately

xver, xhor	Vector specifying nodes to be on the vertical (resp. horizontal) axis of symmetry. The nodes are assumed to flow from left to right. Used by functions <code>force_nodes_on_V_axis()</code> and <code>force_nodes_on_H_axis()</code> respectively, which also move the handles
exact_h, exact_v	Vector specifying nodes to be exactly horizontal or exactly vertical. A node is exactly horizontal (resp. vertical) if the y (resp. x) coordinate of the node is the same as the y (resp. x) coordinate of the handle. Note that the position of an exactly horizontal or vertical node is not restricted, and may be anywhere. Used by functions <code>force_nodes_exactly_horizontal()</code> and <code>force_nodes_exactly_vertical()</code>
symobj	An object representing the symmetry of the knot, usually created by function <code>symmetry_object()</code>
mcdonalds	For vertical symmetry, argument <code>mcdonalds</code> is Boolean, defaulting to <code>FALSE</code> , with <code>TRUE</code> meaning that the symmetric pairs of strands approach the vertical line of symmetry in the same sense (either both moving inward, or both moving outward). It is hard to explain (and named for the gesture one makes when tracing the top two strands a knot with this type of symmetry). The only common knot that needs this is <code>7_2</code>
celtic	Like <code>mcdonalds</code> but for horizontal symmetry
reefknot	Like <code>mcdonalds</code> but for the reefknot
center_crossing	Implements a peculiar type of rotational symmetry in which the strands pass through the geometrical center of the knot projection. The only common knot needing this is <code>9_29</code>
minsymvec	A “minimal symmetric vector”. This is a numeric vector containing just the independent degrees of freedom of a knot, after symmetry constraints have been imposed. The idea is that one may optimize a <code>minsymvec</code> object using <code>nlm()</code> , and then reconstruct a knot using <code>make_minobj_from_minsymvec()</code> together with a symmetry object
vec	A vector, given to function <code>minsymvec()</code>
overunderobj	A matrix specifying the overs and the unders; a two-column matrix with rows corresponding to pairs of strands intersecting. The first element of a row identifies the overstrand and the second element specifies the understrand

## Details

Function `symmetry_object()` creates a symmetry object from `Mver` et seq, but if given a knot object, returns the embedded symmetry object.

There are seven types of symmetry that may be imposed on a knot. These are imposed by the following seven `force_nodes_foo()` functions:

- Functions `force_nodes_mirror_images_LR()` and `force_nodes_mirror_images_UD()` symmetrize a knot about a vertical (resp. horizontal) axis by taking ordered pairs of nodes, specified by matrix `Mver` (resp. `Mhor`) and forcing the second node to be symmetrically placed with respect to the first. It does the same thing to the handles too.



- Functions `force_nodes_exactly_horizontal()` and `force_nodes_exactly_vertical()` force nodes to be exactly horizontal (resp. vertical) by restricting the position of their handles. Nodes so forced do not need to be on an axis of symmetry; they can be anywhere
- Functions `force_nodes_on_V_axis()` and `force_nodes_on_H_axis()` force nodes specified by `xver` (resp. `xhor`) to be on the vertical (resp. horizontal) axis, and to have appropriately placed handles
- Function `force_nodes_rotational()` imposes the rotational symmetry specified by `Mrot`

Function `symmetrize()` imposes the seven kinds of symmetry by calling each of the `force_nodes_foo()` functions in turn.

Function `tag_notneeded()` is an internal function, not really intended for the end-user. It takes a `minobj` object and marks a maximal set of dependent entries with a 'not needed' value. The values of the entries so marked may be determined by a combination of the imposed symmetry relations and the unmarked values. The unmarked entries constitute a `minsymvec` object (see above). These are the *real* degrees of freedom in the symmetrical knot. Only these unmarked values are modified by the optimization routines in `knotoptim()`

### Note

You can achieve up-down symmetry (that is, a horizontal line of symmetry) by making a left-right symmetric knot and rotating by 90 degrees. D'oh.

### Author(s)

Robin K.S. Hankin

### Examples

```
# each row of M = a pair of symmetrical nodes; each element of v is a
# node on the vertical axis

M <- matrix(c(6,4,13,11,7,3,2,8,9,1,14,10),byrow=TRUE,ncol=2)
v <- c(5,12) # on vertical axis

sym_7_3 <- symmetry_object(k7_3, M, v)

k <- symmetrize(as.minobj(k7_3), sym_7_3)

knotplot2(k) #nice and symmetric!

## OK now convert to and from a minimal vector for a symmetrical knot:

mii <- make_minsymvec_from_minobj(k, sym_7_3)
pii <- make_minobj_from_minsymvec(mii,sym_7_3)
knotplot2(pii)

## So 'mii' is a minimal vector for a symmetrical knot, and 'pii' is
```

```
## the corresponding minobj object. Note that you can mess about with
## mii, but whatever you do the resulting knot is still symmetric:

mii[2] <- 1000
knotplot2(make_minobj_from_minsymvec(mii,sym_7_3)) # still symmetric.

## and, in particular, you can optimize the badness, using nlm():

## Not run:
fun <- function(m){badness(make_minobj_from_minsymvec(m,sym_7_3))}
o <- nlm(fun,mii,iterlim=4,print.level=2)

knotplot2(make_minobj_from_minsymvec(o$estimate,sym_7_3))

## End(Not run)
```

---

utilities

*Various utilities for knots*


---

## Description

Various utilities for knots including reading files and creating objects

## Usage

```
controlpoints(x)
inkscape(x)
minobj(x)
knotvec(x)
make_controlpoints_from_ink(a)
make_minobj_from_ink(a)
make_minobj_from_vector(vec)
make_ink_from_minobj(x)
make_inkscape_from_controlpoints(b)
make_minobj_from_knot(k)
make_knotvec_from_minobj(x)
```

## Arguments

x	Suitable object for coercion; see details
a	An inkscape object: a two column matrix with rows representing the positions of nodes and control points
b	A controlpoints object
k	An object of class knot
vec	A vector of reals

## Details

Functions `inkscape()`, `minobj()`, and `knotvec()` are low-level functions; these are the only places that objects have their classes assigned directly. These functions are not user-friendly and require very specific types of object; they perform some checks but are not really intended for the user. Functions `as.foo()` are much more user-friendly, and are documented at `as.Rd`.

Functions `make_foo_from_bar()` coerce `bar` objects into `foo` objects. Functions that involve symmetry are documented at `symmetry.Rd`.

Objects of class `inkscape` are in the form of a two-column matrix, with rows corresponding to 2D positions. The rows correspond to the  $(x, y)$  coordinates of points as held in the `inkscape` file.

There is quite a lot of redundancy in an `inkscape` object:

- The first row of an `inkscape` object is equal to the last row (this follows from the fact that the path is closed).
- If  $n = 0$  modulo 3, then `a[n+2, ] - a[n+1, ] == a[n+1, ] - a[n, ]`, corresponding to the fact that the handles are symmetric in `inkscape`. This is visualised best by `knotplot2(k4_1, ink=TRUE, seg=TRUE)`

Look at functions `make_inkscape_from_minobj()` and `make_minobj_from_ink()` to see this from a symbolic perspective. The vignette also gives some details.

The `minobj` class is a ‘MINimal OBJect’. Objects of class `minobj` are a list of two elements: `$node` and `$handle_A`. Each element has rows corresponding to 2D positions, the same as `inkscape` objects. Element `$node` shows the positions of the nodes, and element `$handle_A` shows the positions of (one of) the handles; the other handle is symmetrically positioned with respect to its node. Use `knotplot2(k4_1, node=TRUE, seg=TRUE)` to see the meaning of the entries; the nodes are indicated by a square and the handles by circles.

NB: objects of class `minobj` have no *redundancy* in the sense that changing any entry of either `$node` or `$handle_A` results in modifying the corresponding `inkscape` diagram. However, doing this to a knot which has imposed symmetry conditions (ie a nontrivial `symobj`) may introduce asymmetry into the `inkscape` diagram. For example, one might take a `minobj` object whose knot diagram is left-right mirror symmetric (eg `k6_2`) and alter one of the handle positions. Then the resulting `inkscape` object will be asymmetric. There is an example using `k6_2` below.

An object of class `controlpoints` is a list of matrices of size 4-by-2. For each matrix, the four rows correspond to the points in 2D Cartesian space needed to specify a Bezier curve; further details and examples are given in `bezier.Rd`. There is lots of redundancy in a `controlpoints` object because the `inkscape` nodes are symmetric nodes with diametrically opposed handles.

The `knotvec` class is a named vector of independent reals suitable for use with optimization routines.

None of the functions here deal with symmetry relations. This is documented at `symmetry.Rd`.

## Author(s)

Robin K. S. Hankin

## See Also

[as,symmetrize](#)

**Examples**

```
a <- as.minobj(k6_3)
plot(a$node,asp=1,pch=16)
segments(x0=a[[1]][,1],y0=a[[1]][,2],x1=a[[2]][,1],y1=a[[2]][,2],
main="handle direction follows the string path")
points(getstringpoints(a),type='l',col='gray',lwd=0.4)
```

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