Sage 9.4 Reference Manual: Knot Theory

Release 9.4

The Sage Development Team

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CHAPTER

ONE

KNOTS

AUTHORS:

- Miguel Angel Marco Buzunariz
- · Amit Jamadagni

class sage.knots.knot.Knot(data, check=True)

Bases: sage.knots.link.Link, sage.structure.element.Element

A knot.

A knot is defined as embedding of the circle \mathbb{S}^1 in the 3-dimensional sphere \mathbb{S}^3 , considered up to ambient isotopy. They represent the physical idea of a knotted rope, but with the particularity that the rope is closed. That is, the ends of the rope are joined.

See also:

Link

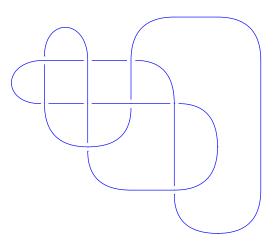
INPUT:

- data see *Link* for the allowable inputs
- check optional, default True. If True, make sure that the data define a knot, not a link

EXAMPLES:

We construct the knot $\mathbf{8}_{14}$ and compute some invariants:

```
sage: B = BraidGroup(4)
sage: K = Knot(B([1,1,1,2,-1,2,-3,2,-3]))
```



```
sage: K.alexander_polynomial()
-2*t^-2 + 8*t^-1 - 11 + 8*t - 2*t^2
sage: K.jones_polynomial()
t^7 - 3*t^6 + 4*t^5 - 5*t^4 + 6*t^3 - 5*t^2 + 4*t + 1/t - 2
sage: K.determinant()
31
sage: K.signature()
-2
```

REFERENCES:

• Wikipedia article Knot_(mathematics)

arf_invariant()

Return the Arf invariant.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: K = Knot(B([-1, 2, 1, 2]))
sage: K.arf_invariant()
0
sage: B = BraidGroup(8)
sage: K = Knot(B([-2, 3, 1, 2, 1, 4]))
sage: K.arf_invariant()
0
sage: K = Knot(B([1, 2, 1, 2]))
sage: K.arf_invariant()
1
```

connected_sum(other)

Return the oriented connected sum of self and other.

Note: We give the knots an orientation based upon the braid representation.

INPUT:

• other - a knot

OUTPUT:

A knot equivalent to the connected sum of self and other.

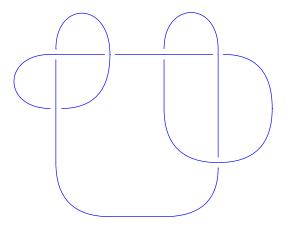
EXAMPLES:

```
sage: B = BraidGroup(2)
sage: trefoil = Knot(B([1,1,1]))
sage: K = trefoil.connected_sum(trefoil); K
Knot represented by 6 crossings
sage: K.braid()
s0^3*s1^-1*s0^3*s1
```

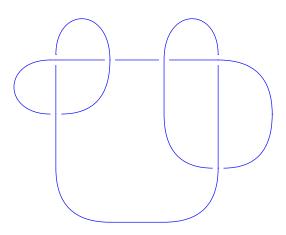
```
sage: rev_trefoil = Knot(B([-1,-1,-1]))
sage: K2 = trefoil.connected_sum(rev_trefoil); K2
Knot represented by 6 crossings
```

(continues on next page)

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```
sage: K2.braid()
s0^3*s1^-1*s0^-3*s1
```



Observe that both knots have according dowker_notation (showing that the constructing from DT-code may not be unique for non prime knots, see from_dowker_code()):

```
sage: K.dowker_notation()
[(4, 1), (2, 5), (6, 3), (10, 7), (8, 11), (12, 9)]
sage: K2.dowker_notation()
[(4, 1), (2, 5), (6, 3), (7, 10), (11, 8), (9, 12)]
sage: K.homfly_polynomial() == K2.homfly_polynomial()
False
```

REFERENCES:

• Wikipedia article Connected_sum

dt_code()

Return the DT code of self.

ALGORITHM:

The DT code is generated by the following way:

Start moving along the knot, as we encounter the crossings we start numbering them, so every crossing has two numbers assigned to it once we have traced the entire knot. Now we take the even number associated with every crossing.

The following sign convention is to be followed:

Take the even number with a negative sign if it is an overcrossing that we are encountering.

OUTPUT: DT code representation of the knot

EXAMPLES:

class sage.knots.knot.Knots

Bases: sage.misc.fast_methods.Singleton, sage.structure.parent.Parent

The set for all knots, as a monoid for the connected sum.

Element

alias of Knot

an_element()

Return the trefoil knot.

EXAMPLES:

```
sage: Knots().an_element()
Knot represented by 3 crossings
```

from_dowker_code(code)

Build a knot from a Dowker-Thistlethwaite code.

The Dowker-Thistlethwaite code of a knot diagram is defined as follows.

Start following the knot diagram at some regular point. Label the crossings by a number (starting from number 1) in the order in which they are met. At the end, every crossing gets numbered twice, once by an even number and once by an odd number. When meeting an over-crossing with even number, use instead the negative of this even number as label.

Then the set of crossings gives a set of pairs (odd, even). Sort this set according to the odd component, and then keep only the even components in the same order. This is the Dowker-Thistlethwaite code.

INPUT:

a list of signed even numbers, the Dowker-Thistlethwaite code of a knot

OUTPUT:

a knot

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Warning: In general the Dowker-Thistlethwaite code does not describe a knot uniquely. It is not only insensitive on mirror images, but may also mix up non prime knots. For example [4, 6, 2, 10, 12, 8] describes the connected sum of two trefoil knots, as well as the connected sum of a trefoil with its mirror (see the corresponding example in the documentation of connected_sum()).

EXAMPLES:

```
sage: W = Knots()
sage: K1 = W.from_dowker_code([8,10,2,12,4,6])
sage: K1.dowker_notation()
[(5, 2), (9, 4), (11, 6), (1, 8), (3, 10), (7, 12)]

sage: W.from_dowker_code([6,10,14,12,16,2,18,4,8])
Knot represented by 9 crossings

sage: W.from_dowker_code([4,8,10,-14,2,-16,-18,-6,-12])
Knot represented by 9 crossings

sage: K3 = W.from_dowker_code([6,-12,2,8,-4,-10]); K3
Knot represented by 6 crossings
sage: K3.dowker_notation()
[(5, 2), (4, 9), (1, 6), (7, 8), (10, 11), (12, 3)]
```

See also:

dowker_notation()

REFERENCES:

- · Wikipedia article Dowker_notation
- http://katlas.org/wiki/DT_(Dowker-Thistlethwaite)_Codes

from_gauss_code(gauss)

Build a knot from a signed Gauss code.

This makes some arbitrary choice of orientation.

INPUT:

· a signed Gauss code

OUTPUT:

a knot

EXAMPLES:

```
sage: W = Knots()
sage: K1 = W.from_gauss_code([2, -1, 3, -2, 1, -3])
sage: K1.alexander_polynomial()
t^-1 - 1 + t
```

$from_table(n, k)$

Return a knot from its index in the Rolfsen table.

INPUT:

• n – the crossing number

• k – a positive integer

OUTPUT:

the knot $K_{n,k}$ in the Rolfsen table

EXAMPLES:

```
sage: K1 = Knots().from_table(6,3); K1
Knot represented by 6 crossings
sage: K1.alexander_polynomial()
t^-2 - 3*t^-1 + 5 - 3*t + t^2

sage: K2 = Knots().from_table(8,4); K2
Knot represented by 9 crossings
sage: K2.determinant()
19
sage: K2.signature()
2

sage: K3 = Knots().from_table(10,56); K3
Knot represented by 11 crossings
sage: K3.jones_polynomial()
t^10 - 3*t^9 + 6*t^8 - 9*t^7 + 10*t^6 - 11*t^5 + 10*t^4 - 7*t^3
+ 5*t^2 - 2*t + 1

sage: K4 = Knots().from_table(10,100)
sage: K4.genus()
4
```

REFERENCES:

• KnotAtlas, http://katlas.math.toronto.edu/wiki/The_Rolfsen_Knot_Table

one()

Return the unit of the monoid.

This is the trivial knot.

EXAMPLES:

```
sage: Knots().one()
Knot represented by 0 crossings
```

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CHAPTER

TWO

LINKS

A knot is defined as embedding of the circle \mathbb{S}^1 in the 3-dimensional sphere \mathbb{S}^3 , considered up to ambient isotopy. They represent the physical idea of a knotted rope, but with the particularity that the rope is closed. That is, the ends of the rope are joined.

A link is an embedding of one or more copies of \mathbb{S}^1 in \mathbb{S}^3 , considered up to ambient isotopy. That is, a link represents the idea of one or more tied ropes. Every knot is a link, but not every link is a knot.

Generically, the projection of a link on ${\bf R}^2$ is a curve with crossings. The crossings are represented to show which strand goes over the other. This curve is called a planar diagram of the link. If we remove the crossings, the resulting connected components are segments. These segments are called the edges of the diagram.

REFERENCES:

- Wikipedia article Knot_(mathematics)
- [Col2013]
- [KnotAtlas]

See also:

There are also tables of link and knot invariants at web-pages KnotInfo and LinkInfo. These can be used inside Sage after installing the optional package database_knotinfo (type sage -i database_knotinfo in a command shell, see knotinfo).

AUTHORS:

- Miguel Angel Marco Buzunariz
- · Amit Jamadagni
- Sebastian Oehms (October 2020, add get_knotinfo() and meth: is_isotopic)

class sage.knots.link.Link(data)

Bases: sage.structure.sage_object.SageObject

A link.

A link is an embedding of one or more copies of \mathbb{S}^1 in \mathbb{S}^3 , considered up to ambient isotopy. That is, a link represents the idea of one or more tied ropes. Every knot is a link, but not every link is a knot.

A link can be created by using one of the conventions mentioned below:

Braid:

• The closure of a braid is a link:

```
sage: B = BraidGroup(8)
sage: L = Link(B([-1, -1, -1, -2, 1, -2, 3, -2, 3]))
sage: L
Link with 1 component represented by 9 crossings
sage: L = Link(B([1, 2, 1, -2, -1]))
sage: L
Link with 2 components represented by 5 crossings
```

Note: The strands of the braid that have no crossings at all are removed.

· Oriented Gauss Code:

Label the crossings from 1 to n (where n is the number of crossings) and start moving along the link. Trace every component of the link, by starting at a particular point on one component of the link and writing down each of the crossings that you encounter until returning to the starting point. The crossings are written with sign depending on whether we cross them as over or undercrossing. Each component is then represented as a list whose elements are the crossing numbers. A second list of +1 and -1's keeps track of the orientation of each crossing:

For links there may be more than one component and the input is as follows:

```
sage: L = Link([[[-1, 2], [-3, 4], [1, 3, -4, -2]], [-1, -1, 1, 1]])
sage: L
Link with 3 components represented by 4 crossings
```

• Planar Diagram (PD) Code:

The diagram of the link is formed by segments that are adjacent to the crossings. Label each one of this segments with a positive number, and for each crossing, write down the four incident segments. The order of these segments is clockwise, starting with the incoming undercrossing.

There is no particular distinction between knots and links for this input.

EXAMPLES:

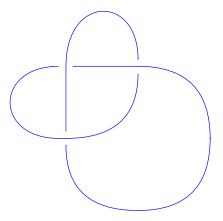
One of the representations of the trefoil knot:

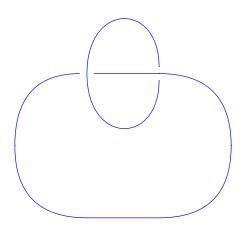
```
sage: L = Link([[1, 5, 2, 4], [5, 3, 6, 2], [3, 1, 4, 6]])
sage: L
Link with 1 component represented by 3 crossings
```

One of the representations of the Hopf link:

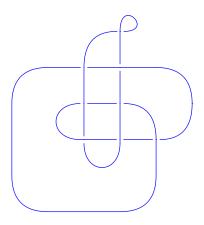
```
sage: L = Link([[1, 4, 2, 3], [4, 1, 3, 2]])
sage: L
Link with 2 components represented by 2 crossings
```

We can construct links from the braid group:

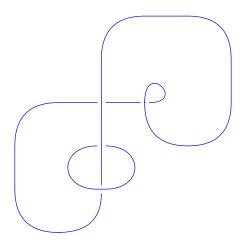




```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, -1, -1, -2, 1, -2, 3, -2]))
sage: L
Link with 2 components represented by 8 crossings
```

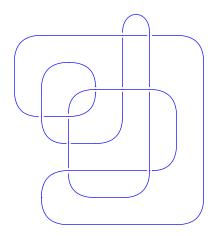


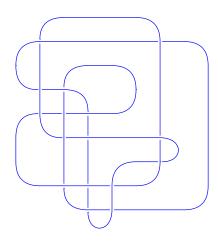
```
sage: L = Link(B([1, 2, 1, 3]))
sage: L
Link with 2 components represented by 4 crossings
```



We construct the "monster" unknot using a planar code, and then construct the oriented Gauss code and braid representation:

We construct the Ochiai unknot by using an oriented Gauss code:

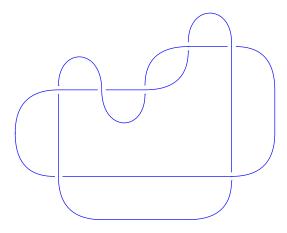




We construct the knot 7_1 and compute some invariants:

```
sage: B = BraidGroup(2)
sage: L = Link(B([1]*7))
```

```
sage: L.alexander_polynomial()
t^-3 - t^-2 + t^-1 - 1 + t - t^2 + t^3
sage: L.jones_polynomial()
-t^10 + t^9 - t^8 + t^7 - t^6 + t^5 + t^3
sage: L.determinant()
7
sage: L.signature()
-6
```



The links here have removed components in which no strand is used:

```
sage: B = BraidGroup(8)
sage: b = B([1])
sage: L = Link(b)
sage: b.components_in_closure()
7
sage: L.number_of_components()
1
sage: L.braid().components_in_closure()
1
sage: L.braid().parent()
Braid group on 2 strands
```

Warning: Equality of knots is done by comparing the corresponding braids, which may give false negatives.

Note: The behavior of removing unused strands from an element of a braid group may change without notice in the future. Do not rely on this feature.

Todo: Implement methods to creating new links from previously created links.

alexander_polynomial(var='t')

Return the Alexander polynomial of self.

INPUT:

• var – (default: 't') the variable in the polynomial

EXAMPLES:

We begin by computing the Alexander polynomial for the figure-eight knot:

```
sage: B = BraidGroup(3)
sage: L = Link(B([1, -2, 1, -2]))

(continues on next page)
```

```
sage: L.alexander_polynomial()
-t^-1 + 3 - t
```

The "monster" unknot:

```
sage: L = Link([[3,1,2,4],[8,9,1,7],[5,6,7,3],[4,18,6,5],
...: [17,19,8,18],[9,10,11,14],[10,12,13,11],
...: [12,19,15,13],[20,16,14,15],[16,20,17,2]])
sage: L.alexander_polynomial()
1
```

Some additional examples:

```
sage: B = BraidGroup(2)
sage: L = Link(B([1]))
sage: L.alexander_polynomial()
1
sage: L = Link(B.one())
sage: L.alexander_polynomial()
1
sage: B = BraidGroup(3)
sage: L = Link(B([1, 2, 1, 2]))
sage: L.alexander_polynomial()
t^-1 - 1 + t
```

When the Seifert surface is disconnected, the Alexander polynomial is defined to be 0:

```
sage: B = BraidGroup(4)
sage: L = Link(B([1,3]))
sage: L.alexander_polynomial()
0
```

arcs(presentation='pd')

Return the arcs of self.

Arcs are the connected components of the planar diagram.

INPUT:

- presentation one of the following:
 - 'pd' the arcs are returned as lists of parts in the PD code
 - 'gauss_code' the arcs are returned as pieces of the Gauss code that start with a negative number, and end with the following negative one; of there exist a closed arc, it is returned as a list of positive numbers only

OUTPUT:

A list of lists representing the arcs based upon presentation.

EXAMPLES:

```
sage: K = Knot([[[1,-2,3,-1,2,-3]],[1,1,1]])
sage: K.arcs()
[[1, 2], [3, 4], [5, 6]]
```

(continues on next page)

```
sage: K.arcs(presentation='gauss_code')
[[-3, 1, -2], [-2, 3, -1], [-1, 2, -3]]
```

```
sage: L = Link([[1, 2, 3, 4], [3, 2, 1, 4]])
sage: L.arcs()
[[2, 4], [1], [3]]
sage: L.arcs(presentation='gauss_code')
[[-2, -1], [-1, -2], [2, 1]]
sage: L.gauss_code()
[[-1, -2], [2, 1]]
```

braid()

Return a braid representation of self.

OUTPUT: an element in the braid group

Warning: For the unknot with no crossings, this returns the identity of the braid group with 2 strands because this disregards strands with no crossings.

EXAMPLES:

```
sage: L = Link([[2, 3, 1, 4], [4, 1, 3, 2]])
sage: L.braid()
s^2
sage: L = Link([[[-1, 2, -3, 1, -2, 3]], [-1, -1, -1]])
sage: L.braid()
s^-3
sage: L = Link([[1,8,2,7], [8,4,9,5], [3,9,4,10], [10,1,7,6], [5,3,6,2]])
sage: L.braid()
(s0*s1^-1)^2*s1^-1
```

colorings(n)

Return the n-colorings of self.

INPUT:

• n – the number of colors to consider

If n is not a prime number, it is replaced by the smallest prime number that is larger than n.

OUTPUT:

a list with the colorings. Each coloring is represented as a dictionary that maps a tuple of the edges forming each arc (as in the PD code) to the index of the corresponding color.

EXAMPLES:

(continues on next page)

```
{(1, 2): 2, (3, 4): 1, (5, 6): 0}]

sage: K.pd_code()

[[4, 1, 5, 2], [2, 5, 3, 6], [6, 3, 1, 4]]

sage: K.arcs('pd')

[[1, 2], [3, 4], [5, 6]]
```

REFERENCES:

- Wikipedia article Fox_n-coloring
- Chapter 3 of [Liv1993]

See also:

is_colorable()

determinant()

Return the determinant of self.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, 2, 1, 2]))
sage: L.determinant()
1
sage: B = BraidGroup(8)
sage: L = Link(B([2, 4, 2, 3, 1, 2]))
sage: L.determinant()
3
sage: L = Link(B([1]*16 + [2,1,2,1,2,2,2,2,2,2,2,1,2,1,2,-1,2,-2]))
sage: L.determinant()
65
```

dowker_notation()

Return the Dowker notation of self.

Similar to the PD code we number the components, so every crossing is represented by four numbers. We focus on the incoming entities of the under and the overcrossing. It is the pair of incoming undercrossing and the incoming overcrossing. This information at every crossing gives the Dowker notation.

OUTPUT:

A list containing the pair of incoming under cross and the incoming over cross.

EXAMPLES:

fundamental_group(presentation='wirtinger')

Return the fundamental group of the complement of self.

INPUT:

- presentation string; one of the following:
 - 'wirtinger' (default) the Wirtinger presentation (see Wikipedia article Link_group)
 - 'braid' the presentation is given by the braid action on the free group (see chapter 2 of [Bir1975])

OUTPUT:

· a finitely presented group

EXAMPLES:

```
sage: L = Link([[1, 2, 3, 4], [3, 2, 1, 4]])
sage: L.fundamental_group()
Finitely presented group < x0, x1, x2 | x1*x0^-1*x2^-1*x0, x2*x0*x1^-1*x0^-1 >
sage: L.fundamental_group('braid')
Finitely presented group < x0, x1 | 1, 1 >
```

We can see, for instance, that the two presentations of the group of the figure eight knot correspond to isomorphic groups:

```
sage: K8 = Knot([[[1, -2, 4, -3, 2, -1, 3, -4]], [1, 1, -1, -1]])
sage: GA = K8.fundamental_group()
sage: GA
Finitely presented group < x0, x1, x2, x3 |
  x2*x0*x3^{-1}*x0^{-1}, x0*x2*x1^{-1}*x2^{-1},
  x1*x3^{-1}x2^{-1}x3, x3*x1^{-1}x0^{-1}x1 >
sage: GB = K8.fundamental_group(presentation='braid')
sage: GB
Finitely presented group < x0, x1, x2 | x1*x2^-1*x1^-1*x0*x1*x2*x1*x2^-1*x1^-
 \rightarrow 1*x0^{-1}x1^{x}2^{x}1^{-1}x0^{-1}, x1^{x}x2^{-1}x1^{-1}x0^{x}1^{x}x2^{x}1^{-1}x2^{-1}x1^{-1}x0^{x}
 \rightarrow 1*x1*x2*x1^{-1}x0*x1*x2*x1*x2^{-1}x1^{-1}x0^{-1}x1^{-1}x0^{-1}x1^{-1}x0^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1}x1^{-1
 \rightarrow 1*x0*x1*x2*x1^-1*x2^-1 >
sage: GA.simplified()
Finitely presented group < x0, x1 \mid
  x1^{-1}x0^{x}1^{x}0^{-1}x1^{x}1^{x}0^{x}1^{-1}x0^{-1}x1^{x}0^{-1} > 0
sage: GB.simplified()
Finitely presented group < x0, x2 |
  x2^{-1}x0^{x}2^{-1}x0^{x}2^{-1}x0^{-1}x2^{x}0^{x}2^{-1}x0^{x}2^{x}0^{-1} >
```

gauss_code()

Return the Gauss code of self.

The Gauss code is generated by the following procedure:

- a. Number the crossings from 1 to n.
- b. Select a point on the knot and start moving along the component.
- c. At each crossing, take the number of the crossing, along with sign, which is if it is an undercrossing and + if it is a overcrossing.

EXAMPLES:

```
sage: L = Link([[1, 4, 2, 3], [4, 1, 3, 2]])
sage: L.gauss_code()
[[-1, 2], [1, -2]]
sage: B = BraidGroup(8)
sage: L = Link(B([1, -2, 1, -2, -2]))
sage: L.gauss_code()
[[-1, 3, -4, 5], [1, -2, 4, -5, 2, -3]]
sage: L = Link([[[-1, 2], [-3, 4], [1, 3, -4, -2]], [-1, -1, 1, 1]])
sage: L.gauss_code()
[[-1, 2], [-3, 4], [1, 3, -4, -2]]
```

genus()

Return the genus of self.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, 3, 1, 3]))
sage: L.genus()
0
sage: L = Link(B([1,3]))
sage: L.genus()
0
sage: B = BraidGroup(8)
sage: L = Link(B([-2, 4, 1, 6, 1, 4]))
sage: L.genus()
0
sage: L = Link(B([1, 2, 1, 2]))
sage: L = Link(B([1, 2, 1, 2]))
sage: L.genus()
1
```

get_knotinfo(mirror_version=True, unique=True)

Identify this link as an item of the KnotInfo database (if possible).

INPUT:

- mirror_version boolean (default is True). If set to False the result of the method will be just the instance of *KnotInfoBase* (by default the result is a tuple of the instance and a boolean, see explanation of the output below)
- unique boolean (default is True). This only affects the case where a unique identification is not possible. If set to False you can obtain a matching list (see explanation of the output below)

OUTPUT:

A tuple (K, m) where K is an instance of *KnotInfoBase* and m a boolean (for chiral links) telling if self corresponds to the mirrored version of K or not. The value of m is None for amphicheiral links and ? if it cannot be determined uniquely and the keyword option unique=False is given.

For proper links, if the orientation mutant cannot be uniquely determined, K will be a series of links gathering all links having the same unoriented name, that is an instance of *KnotInfoSeries*.

If mirror_version is set to False then the result is just K (that is: m is suppressed).

If it is not possible to determine a unique result a NotImplementedError will be raised. To avoid this you can set unique to False. You will get a list of matching candidates instead.

Note: The identification of proper links may fail to be unique due to the following fact: In opposite to

the database for knots, there are pairs of oriented mutants of an unoriented link which are isotopic to each other. For example L5a1_0 and L5a1_1 is such a pair.

This is because all combinatorial possible oriented mutants are listed with individual names regardless whether they are pairwise non isotopic or not. In such a case the identification is not unique and therefore a series of the links will be returned which gathers all having the same unoriented name.

To obtain the individual oriented links being isotopic to self use the keyword unique (see the examples for L2a1_1 and L5a1_0 below).

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = Link([[4,2,5,1], [10,3,11,4], [5,16,6,17], [7,12,8,13],
. . . . :
                [18,9,19,10], [2,11,3,12], [13,20,14,21], [15,6,16,7],
                [22,18,1,17], [8,19,9,20], [21,14,22,15]])
. . . . :
sage: L.get_knotinfo()
                                  # optional - database_knotinfo
(<KnotInfo.K11n_121: '11n_121'>, True)
sage: K = KnotInfo.K10_25
                                  # optional - database_knotinfo
sage: 1 = K.link()
                                  # optional - database_knotinfo
                                  # optional - database_knotinfo
sage: 1.get_knotinfo()
(<KnotInfo.K10_25: '10_25'>, False)
```

Knots with more than 12 and proper links having more than 11 crossings cannot be identified. In addition non prime links or even links whose HOMFLY-PT polynomial is not irreducible cannot be identified:

Lets identify the monster unknot:

Usage of option mirror_version:

```
sage: L.get_knotinfo(mirror_version=False) == KnotInfo.K0_1
True
```

Usage of option unique:

```
sage: 1 = K.link(K.items.gauss_notation) # optional - database_knotinfo
sage: 1.get_knotinfo()
                                          # optional - database_knotinfo
Traceback (most recent call last):
NotImplementedError: this link cannot be uniquely determined
use keyword argument `unique` to obtain more details
sage: 1.get_knotinfo(unique=False)
                                          # optional - database_knotinfo
[(<KnotInfo.K10_25: '10_25'>, False), (<KnotInfo.K10_56: '10_56'>, False)]
sage: k11 = KnotInfo.K11n_82.link()
                                          # optional - database_knotinfo
sage: k11m = k11.mirror_image()
                                          # optional - database_knotinfo
sage: k11m.braid()
                                          # optional - database_knotinfo
s0*s1^-1*s0*s2^-1*s1*(s1*s2^-1)^2*s2^-2
sage: k11m.get_knotinfo()
                                          # optional - database_knotinfo
Traceback (most recent call last):
NotImplementedError: mirror type of this link cannot be uniquely determined
use keyword argument `unique` to obtain more details
sage: k11m.get_knotinfo(unique=False)
                                          # optional - database_knotinfo
[(<KnotInfo.K11n_82: '11n_82'>, '?')]
sage: t = (-1, 2, -1, 2, -1, 3, -2, 3, -2)
sage: 19 = Link(BraidGroup(4)(t))
sage: 19.get_knotinfo()
                                          # optional - database_knotinfo
Traceback (most recent call last):
NotImplementedError: this link cannot be uniquely determined
use keyword argument `unique` to obtain more details
sage: 19.get_knotinfo(unique=False)
                                          # optional - database_knotinfo
[(<KnotInfo.L9n25_0_0: 'L9n25{0,0}'>, False),
(<KnotInfo.L9n25_1_1: 'L9n25{1,1}'>, False)]
sage: t = (1, 2, 3, -4, 3, -2, -1, 3, -2, 3, -2, 3, -4, 3, -2)
sage: 115 = Link(BraidGroup(5)(t))
sage: l15.get_knotinfo()
                                          # optional - database_knotinfo
Traceback (most recent call last):
NotImplementedError: this link having more than 11 crossings cannot be uniquely.
→determined
use keyword argument `unique` to obtain more details
sage:l15.get_knotinfo(unique=False)
                                          # optional - database_knotinfo
[(<KnotInfo.L11a1_0: 'L11a1{0}'>, False),
(<KnotInfo.L11a1_1: 'L11a1{1}'>, False)]
```

Furthermore, if the result is a complete series of oriented links having the same unoriented name (according to the note above) the option can be used to achieve more detailed information:

```
sage: L2a1 = Link(b**2)
sage: L2a1.get_knotinfo()
(Series of links L2a1, None)
sage: L2a1.get_knotinfo(unique=False)
[(<KnotInfo.L2a1_0: 'L2a1{0}'>, True), (<KnotInfo.L2a1_1: 'L2a1{1}'>, False)]
sage: KnotInfo.L5a1_0.inject()
Defining L5a1_0
sage: 15 = Link(L5a1_0.braid())
sage: 15.get_knotinfo()
(Series of links L5a1, False)
sage: _[0].inject()
Defining L5a1
sage: list(L5a1)
[<KnotInfo.L5a1_0: 'L5a1{0}'>, <KnotInfo.L5a1_1: 'L5a1{1}'>]
sage: 15.get_knotinfo(unique=False)
[(<KnotInfo.L5a1_0: 'L5a1{0}'>, False), (<KnotInfo.L5a1_1: 'L5a1{1}'>, False)]
```

Clarifying the series around the Perko pair (Wikipedia article Perko pair):

```
sage: for i in range(160, 166):  # optional - database_knotinfo
....:  K = Knots().from_table(10, i)
....:  print('%s_%s' %(10, i), '--->', K.get_knotinfo())
10_160 ---> (<KnotInfo.K10_160: '10_160'>, False)
10_161 ---> (<KnotInfo.K10_161: '10_161'>, True)
10_162 ---> (<KnotInfo.K10_162: '10_162'>, False)
10_163 ---> (<KnotInfo.K10_163: '10_163'>, False)
10_164 ---> (<KnotInfo.K10_164: '10_164'>, False)
10_165 ---> (<KnotInfo.K10_165: '10_165'>, True)
```

Clarifying ther Perko series against SnapPy:

```
sage: import snappy
                                       # optional - snappy
Plink failed to import tkinter.
sage: from sage.knots.knotinfo import KnotInfoSeries
sage: KnotInfoSeries(10, True, True) # optional - database_knotinfo
Series of knots K10
                                     # optional - database_knotinfo
sage: _.inject()
Defining K10
sage: for i in range(160, 166):
                                      # optional - database_knotinfo snappy
. . . . .
         K = K10(i)
         k = K.link(K.items.name, snappy=True)
         print(k, '--->', k.sage_link().get_knotinfo())
<Link 10_160: 1 comp; 10 cross> ---> (<KnotInfo.K10_160: '10_160'>, False)
<Link 10_161: 1 comp; 10 cross> ---> (<KnotInfo.K10_161: '10_161'>, True)
<Link 10_162: 1 comp; 10 cross> ---> (<KnotInfo.K10_161: '10_161'>, False)
<Link 10_163: 1 comp; 10 cross> ---> (<KnotInfo.K10_162: '10_162'>, False)
<Link 10_164: 1 comp; 10 cross> ---> (<KnotInfo.K10_163: '10_163'>, False)
<Link 10_165: 1 comp; 10 cross> ---> (<KnotInfo.K10_164: '10_164'>, False)
sage: snappy.Link('10_166')
                                       # optional - snappy
<Link 10_166: 1 comp; 10 cross>
```

(continues on next page)

```
sage: _.sage_link().get_knotinfo() # optional - database_knotinfo snappy
(<KnotInfo.K10_165: '10_165'>, True)
```

Another pair of confusion (see the corresponding Warning):

```
sage: Ks10_86 = snappy.Link('10_86')  # optional - snappy
sage: Ks10_83 = snappy.Link('10_83')  # optional - snappy
sage: Ks10_86.sage_link().get_knotinfo() # optional - snappy
(<KnotInfo.K10_83: '10_83'>, True)
sage: Ks10_83.sage_link().get_knotinfo() # optional - snappy
(<KnotInfo.K10_86: '10_86'>, False)
```

homfly_polynomial(var1=None, var2=None, normalization='lm')

Return the HOMFLY polynomial of self.

The HOMFLY polynomial P(K) of a link K is a Laurent polynomial in two variables defined using skein relations and for the unknot U, we have P(U) = 1.

INPUT:

- var1 (default: 'L') the first variable. If normalization is set to az resp. vz the default is a resp. v
- var2 (default: 'M') the second variable. If normalization is set to az resp. vz the default is z
- normalization (default: 1m) the system of coordinates and can be one of the following:
 - 'lm' corresponding to the Skein relation $L \cdot P(K_+) + L^{-1} \cdot P(K_-) + M \cdot P(K_0) = 0$
 - 'az' corresponding to the Skein relation $a \cdot P(K_+) a^{-1} \cdot P(K_-) = z \cdot P(K_0)$
 - 'vz' corresponding to the Skein relation $v^{-1} \cdot P(K_+) v \cdot P(K_-) = z \cdot P(K_0)$

where $P(K_+)$, $P(K_-)$ and $P(K_0)$ represent the HOMFLY polynomials of three links that vary only in one crossing; that is the positive, negative, or smoothed links respectively

OUTPUT:

A Laurent polynomial over the integers.

Note: Use the 'az' normalization to agree with the data in [KnotAtlas]

Use the 'vz' normalization to agree with the data KnotInfo.

EXAMPLES:

We give some examples:

```
sage: g = BraidGroup(2).gen(0)
sage: K = Knot(g^5)
sage: K.homfly_polynomial()
L^-4*M^4 - 4*L^-4*M^2 + 3*L^-4 - L^-6*M^2 + 2*L^-6
```

The Hopf link:

```
sage: L = Link([[1,3,2,4],[4,2,3,1]])
sage: L.homfly_polynomial('x', 'y')
-x^-1*y + x^-1*y^-1 + x^-3*y^-1
```

Another version of the Hopf link where the orientation has been changed. Therefore we substitute $x \mapsto L^{-1}$ and $y \mapsto M$:

```
sage: L = Link([[1,4,2,3], [4,1,3,2]])
sage: L.homfly_polynomial()
L^3*M^-1 - L*M + L*M^-1
sage: L = Link([[1,4,2,3], [4,1,3,2]])
sage: L.homfly_polynomial(normalization='az')
a^3*z^-1 - a*z - a*z^-1
```

The figure-eight knot:

```
sage: L = Link([[2,1,4,5], [5,6,7,3], [6,4,1,9], [9,2,3,7]])
sage: L.homfly_polynomial()
-L^2 + M^2 - 1 - L^-2
sage: L.homfly_polynomial('a', 'z', 'az')
a^2 - z^2 - 1 + a^-2
```

The "monster" unknot:

Comparison with KnotInfo:

```
sage: KI, m = K.get_knotinfo(); KI, m
  (<KnotInfo.K5_1: '5_1'>, False)
sage: K.homfly_polynomial(normalization='vz') == KI.homfly_polynomial()
True
```

The knot 96:

```
sage: B = BraidGroup(3)
sage: K = Knot(B([-1,-1,-1,-1,-1,-2,1,-2,-2]))
sage: K.homfly_polynomial()
L^10*M^4 - L^8*M^6 - 3*L^10*M^2 + 4*L^8*M^4 + L^6*M^6 + L^10
- 3*L^8*M^2 - 5*L^6*M^4 - L^8 + 7*L^6*M^2 - 3*L^6
sage: K.homfly_polynomial('a', 'z', normalization='az')
-a^10*z^4 + a^8*z^6 - 3*a^10*z^2 + 4*a^8*z^4 + a^6*z^6 - a^10
+ 3*a^8*z^2 + 5*a^6*z^4 - a^8 + 7*a^6*z^2 + 3*a^6
```

REFERENCES:

- Wikipedia article HOMFLY_polynomial
- http://mathworld.wolfram.com/HOMFLYPolynomial.html

is_alternating()

Return whether the given knot diagram is alternating.

Alternating diagram implies every overcross is followed by an undercross or the vice-versa.

We look at the Gauss code if the sign is alternating, True is returned else the knot is not alternating False is returned.

Warning: This does not check if a knot admits an alternating diagram or not. Thus, this term is used differently than in some of the literature, such as in Hoste-Thistlethwaite table.

Note: Links with more than one component are considered to not be alternating (knots) even when such a diagram exists.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, -1, -1, -1]))
sage: L.is_alternating()
False
sage: L = Link(B([1, -2, -1, 2]))
sage: L.is_alternating()
False
sage: L = Link(B([-1, 3, 1, 3, 2]))
sage: L.is_alternating()
False
sage: L = Link(B([1]*16 + [2,1,2,1,2,2,2,2,2,2,1,2,1,2,-1,2,-2]))
sage: L.is_alternating()
False
sage: L = Link(B([-1,2,-1,2]))
sage: L.is_alternating()
True
```

We give the 5_2 knot with an alternating diagram and a non-alternating diagram:

is_colorable(n)

Return whether the link is n-colorable.

A link is n-colorable if its arcs can be painted with n colours, labeled from 0 to n-1, in such a way that at any crossing, the average of the indices of the undercrossings equals twice the index of the overcrossing.

INPUT:

• n – the number of colors to consider

If n is not a prime number, it is replaced by the smallest prime number that is larger than n.

EXAMPLES:

We show that the trefoil knot is 3-colorable:

```
sage: K = Link([[[1, -2, 3, -1, 2, -3]], [1, 1, 1]])
sage: K.is_colorable(3)
True
```

But the figure eight knot is not:

```
sage: K8 = Link([[[1, -2, 4, -3, 2, -1, 3, -4]], [1, 1, -1, -1]])
sage: K8.is_colorable(3)
False
```

REFERENCES:

- Wikipedia article Fox_n-coloring
- Chapter 3 of [Liv1993]

See also:

colorings()

is_isotopic(other)

Check whether self is isotopic to other.

INPUT:

• other – another instance of Link

EXAMPLES:

```
sage: 11 = Link([[2, 9, 3, 10], [4, 13, 5, 14], [6, 11, 7, 12],
                 [8, 1, 9, 2], [10, 7, 11, 8], [12, 5, 13, 6],
                 [14, 3, 1, 4]])
sage: 12 = Link([[1, 8, 2, 9], [9, 2, 10, 3], [3, 14, 4, 1],
                 [13, 4, 14, 5], [5, 12, 6, 13], [11, 6, 12, 7],
                 [7, 10, 8, 11]])
sage: l1.is_isotopic(l2)
True
sage: 13 = 12.mirror_image()
sage: l1.is_isotopic(l3)
False
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L7a7_0_0
                                       # optional - database_knotinfo
sage: L.series(oriented=True).inject() # optional - database_knotinfo
Defining L7a7
sage: L == L7a7(0)
                                        # optional - database_knotinfo
True
sage: 1 = L.link()
                                        # optional - database_knotinfo
                                        # optional - database_knotinfo
sage: l.is_isotopic(L7a7(1).link())
Traceback (most recent call last):
NotImplementedError: comparison not possible!
sage: l.is_isotopic(L7a7(2).link())
                                       # optional - database_knotinfo
sage: 1.is_isotopic(L7a7(3).link()) # optional - database_knotinfo
False
```

is_knot()

Return True if self is a knot.

Every knot is a link but the converse is not true.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([1, 3, 1, -3]))
sage: L.is_knot()
False
sage: B = BraidGroup(8)
sage: L = Link(B([1, 2, 3, 4, 5, 6]))
sage: L.is_knot()
True
```

 ${\tt jones_polynomial} (\textit{variab} = \textit{None}, \textit{skein_normalization} = \textit{False}, \textit{algorithm} = \textit{'jonesrep'})$

Return the Jones polynomial of self.

The normalization is so that the unknot has Jones polynomial 1. If skein_normalization is True, the variable of the result is replaced by a itself to the power of 4, so that the result agrees with the conventions of [Lic1997] (which in particular differs slightly from the conventions used otherwise in this class), had one used the conventional Kauffman bracket variable notation directly.

If variab is None return a polynomial in the variable A or t, depending on the value skein_normalization. In particular, if skein_normalization is False, return the result in terms of the variable t, also used in [Lic1997].

ALGORITHM:

The calculation goes through one of two possible algorithms, depending on the value of algorithm. Possible values are 'jonesrep' which uses the Jones representation of a braid representation of self to compute the polynomial of the trace closure of the braid, and statesum which recursively computes the Kauffman bracket of self. Depending on how the link is given, there might be significant time gains in using one over the other. When the trace closure of the braid is self, the algorithms give the same result.

INPUT:

- variab variable (default: None); the variable in the resulting polynomial; if unspecified, use either a default variable in $\mathbf{Z}[A, A^{-1}]$ or the variable t in the symbolic ring
- skein_normalization boolean (default: False); determines the variable of the resulting polynomial
- algorithm string (default: 'jonesrep'); algorithm to use and can be one of the following:
 - 'jonesrep' use the Jones representation of the braid representation
 - 'statesum' recursively computes the Kauffman bracket

OUTPUT:

If skein_normalization if False, this returns an element in the symbolic ring as the Jones polynomial of the link might have fractional powers when the link is not a knot. Otherwise the result is a Laurent polynomial in variab.

EXAMPLES:

The unknot:

```
sage: B = BraidGroup(9)
sage: b = B([1, 2, 3, 4, 5, 6, 7, 8])
sage: Link(b).jones_polynomial()
1
```

The "monster" unknot:

The Ochiai unknot:

Two unlinked unknots:

```
sage: B = BraidGroup(4)
sage: b = B([1, 3])
sage: Link(b).jones_polynomial()
-sqrt(t) - 1/sqrt(t)
```

The Hopf link:

```
sage: B = BraidGroup(2)
sage: b = B([-1,-1])
sage: Link(b).jones_polynomial()
-1/sqrt(t) - 1/t^(5/2)
```

Different representations of the trefoil and one of its mirror:

```
sage: B = BraidGroup(2)
sage: b = B([-1, -1, -1])
sage: Link(b).jones_polynomial(skein_normalization=True)
-A^{-16} + A^{-12} + A^{-4}
sage: Link(b).jones_polynomial()
1/t + 1/t^3 - 1/t^4
sage: B = BraidGroup(3)
sage: b = B([-1, -2, -1, -2])
sage: Link(b).jones_polynomial(skein_normalization=True)
-A^{-16} + A^{-12} + A^{-4}
sage: R.<x> = LaurentPolynomialRing(GF(2))
sage: Link(b).jones_polynomial(skein_normalization=True, variab=x)
x^{-16} + x^{-12} + x^{-4}
sage: B = BraidGroup(3)
sage: b = B([1, 2, 1, 2])
sage: Link(b).jones_polynomial(skein_normalization=True)
A^4 + A^12 - A^16
```

K11n42 (the mirror of the "Kinoshita-Terasaka" knot) and K11n34 (the mirror of the "Conway" knot) in [KnotAtlas]:

```
sage: B = BraidGroup(4)
sage: K11n42 = Link(B([1, -2, 3, -2, 3, -2, -2, -1, 2, -3, -3, 2, 2]))
sage: K11n34 = Link(B([1, 1, 2, -3, 2, -3, 1, -2, -2, -3, -3]))
sage: bool(K11n42.jones_polynomial() == K11n34.jones_polynomial())
True
```

The two algorithms for computation give the same result when the trace closure of the braid representation is the link itself:

When we have thrown away unknots so that the trace closure of the braid is not necessarily the link itself, this is only true up to a power of the Jones polynomial of the unknot:

```
sage: B = BraidGroup(3)
sage: b = B([1])
sage: L = Link(b)
sage: b.components_in_closure()
2
sage: L.number_of_components()
1
sage: b.jones_polynomial()
-sqrt(t) - 1/sqrt(t)
sage: L.jones_polynomial()
1
sage: L.jones_polynomial()
1
```

khovanov_homology(ring=Integer Ring, height=None, degree=None)

Return the Khovanov homology of the link.

INPUT:

- ring (default: ZZ) the coefficient ring
- height the height of the homology to compute, if not specified, all the heights are computed
- degree the degree of the homology to compute, if not specified, all the degrees are computed

OUTPUT:

The Khovanov homology of the Link. It is given as a dictionary whose keys are the different heights. For each height, the homology is given as another dictionary whose keys are the degrees.

EXAMPLES:

```
sage: K = Link([[[1, -2, 3, -1, 2, -3]],[-1, -1, -1]])
sage: K.khovanov_homology()
{-9: {-3: Z},
```

(continues on next page)

```
-7: {-3: 0, -2: C2},

-5: {-3: 0, -2: Z, -1: 0, 0: 0},

-3: {-3: 0, -2: 0, -1: 0, 0: Z},

-1: {0: Z}}
```

The figure eight knot:

```
sage: L = Link([[1, 6, 2, 7], [5, 2, 6, 3], [3, 1, 4, 8], [7, 5, 8, 4]])
sage: L.khovanov_homology(height=-1)
{-1: {-2: 0, -1: Z, 0: Z, 1: 0, 2: 0}}
```

The Hopf link:

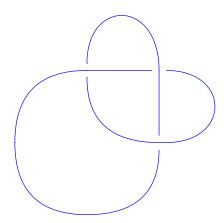
```
sage: B = BraidGroup(2)
sage: b = B([1, 1])
sage: K = Link(b)
sage: K.khovanov_homology(degree = 2)
{2: {2: 0}, 4: {2: Z}, 6: {2: Z}}
```

mirror_image()

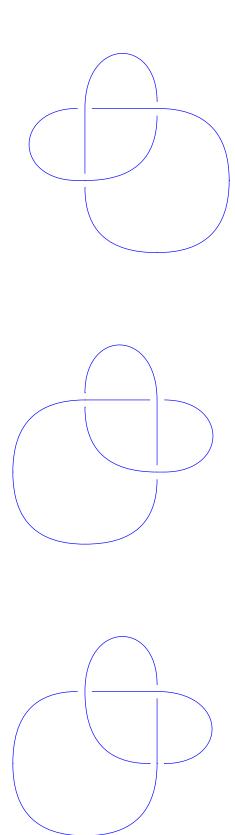
Return the mirror image of self.

EXAMPLES:

```
sage: g = BraidGroup(2).gen(0)
sage: K = Link(g^3)
sage: K2 = K.mirror_image(); K2
Link with 1 component represented by 3 crossings
sage: K2.braid()
s^-3
```



```
sage: K = Knot([[[1, -2, 3, -1, 2, -3]], [1, 1, 1]])
sage: K2 = K.mirror_image(); K2
Knot represented by 3 crossings
sage: K.pd_code()
[[4, 1, 5, 2], [2, 5, 3, 6], [6, 3, 1, 4]]
sage: K2.pd_code()
[[4, 2, 5, 1], [2, 6, 3, 5], [6, 4, 1, 3]]
```



number_of_components()

Return the number of connected components of self.

OUTPUT: number of connected components

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, 3, 1, 3]))
sage: L.number_of_components()
4
sage: B = BraidGroup(8)
sage: L = Link(B([-2, 4, 1, 6, 1, 4]))
sage: L.number_of_components()
5
sage: L = Link(B([1, 2, 1, 2]))
sage: L.number_of_components()
1
sage: L = Link(B.one())
sage: L.number_of_components()
```

omega_signature(omega)

Compute the ω -signature of self.

INPUT:

• ω – a complex number of modulus 1. This is assumed to be coercible to QQbar.

This is defined as the signature of the Hermitian matrix

$$(1-\omega)V + (1-\omega^{-1})V^t$$
,

where V is the Seifert matrix, as explained on page 122 of [Liv1993].

According to [Con2018], this is also known as the Levine-Tristram signature, the equivariant signature or the Tristram-Levine signature.

See also:

```
signature(), seifert_matrix()
```

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: K = Knot(B([1,1,1,2,-1,2,-3,2,-3]))
sage: omega = QQbar.zeta(3)
sage: K.omega_signature(omega)
-2
```

orientation()

Return the orientation of the crossings of the link diagram of self.

EXAMPLES:

(continues on next page)

oriented_gauss_code()

Return the oriented Gauss code of self.

The oriented Gauss code has two parts:

- a. the Gauss code
- b. the orientation of each crossing

The following orientation was taken into consideration for construction of knots:

From the outgoing of the overcrossing if we move in the clockwise direction to reach the outgoing of the undercrossing then we label that crossing as -1.

From the outgoing of the overcrossing if we move in the anticlockwise direction to reach the outgoing of the undercrossing then we label that crossing as +1.

One more consideration we take in while constructing the orientation is the order of the orientation is same as the ordering of the crossings in the Gauss code.

Note: Convention: under is denoted by -1, and over by +1 in the crossing info.

EXAMPLES:

pd_code()

Return the planar diagram code of self.

The planar diagram is returned in the following format.

We construct the crossing by starting with the entering component of the undercrossing, move in the clockwise direction and then generate the list. If the crossing is given by [a, b, c, d], then we interpret this information as:

- 1. a is the entering component of the undercrossing;
- 2. b, d are the components of the overcrossing;

3. c is the leaving component of the undercrossing.

EXAMPLES:

```
sage: L = Link([[[1, -2, 3, -4, 2, -1, 4, -3]], [1, 1, -1, -1]])
sage: L.pd_code()
[[6, 1, 7, 2], [2, 5, 3, 6], [8, 4, 1, 3], [4, 8, 5, 7]]
sage: B = BraidGroup(2)
sage: b = B([1, 1, 1, 1, 1])
sage: L = Link(b)
sage: L.pd_code()
[[2, 1, 3, 4], [4, 3, 5, 6], [6, 5, 7, 8], [8, 7, 9, 10], [10, 9, 1, 2]]
sage: L = Link([[[2, -1], [1, -2]], [1, 1]])
sage: L.pd_code()
[[2, 3, 1, 4], [4, 1, 3, 2]]
sage: L = Link([[1, 2, 3, 3], [2, 4, 5, 5], [4, 1, 7, 7]])
sage: L.pd_code()
[[1, 2, 3, 3], [2, 4, 5, 5], [4, 1, 7, 7]]
```

plot(gap=0.1, component_gap=0.5, solver=None, color='blue', **kwargs)
Plot self.

INPUT:

- gap (default: 0.1) the size of the blank gap left for the crossings
- component_gap (default: 0.5) the gap between isolated components
- solver the linear solver to use, see MixedIntegerLinearProgram.
- color (default: 'blue') a color or a coloring (as returned by *colorings(*).

The usual keywords for plots can be used here too.

EXAMPLES:

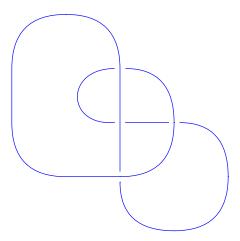
We construct the simplest version of the unknot:

```
sage: L = Link([[2, 1, 1, 2]])
sage: L.plot()
Graphics object consisting of ... graphics primitives
```

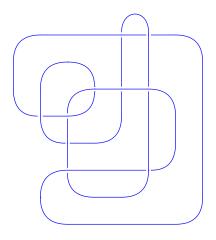


We construct a more interesting example of the unknot:

```
sage: L = Link([[2, 1, 4, 5], [3, 5, 6, 7], [4, 1, 9, 6], [9, 2, 3, 7]])
sage: L.plot()
Graphics object consisting of ... graphics primitives
```

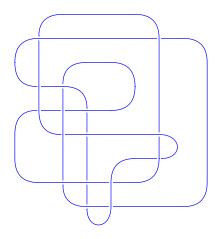


The "monster" unknot:

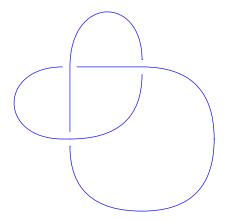


The Ochiai unknot:

One of the representations of the trefoil knot:



```
sage: L = Link([[1, 5, 2, 4], [5, 3, 6, 2], [3, 1, 4, 6]])
sage: L.plot()
Graphics object consisting of 14 graphics primitives
```



The figure-eight knot:

```
sage: L = Link([[2, 1, 4, 5], [5, 6, 7, 3], [6, 4, 1, 9], [9, 2, 3, 7]])
sage: L.plot()
Graphics object consisting of ... graphics primitives
```

The knot K11n121 in [KnotAtlas]:

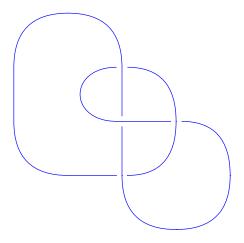
```
sage: L = Link([[4,2,5,1], [10,3,11,4], [5,16,6,17], [7,12,8,13],
....: [18,9,19,10], [2,11,3,12], [13,20,14,21], [15,6,16,7],
....: [22,18,1,17], [8,19,9,20], [21,14,22,15]])
sage: L.plot()
Graphics object consisting of ... graphics primitives
```

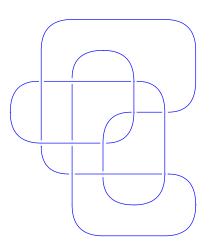
One of the representations of the Hopf link:

```
sage: L = Link([[1, 4, 2, 3], [4, 1, 3, 2]])
sage: L.plot()
```

(continues on next page)

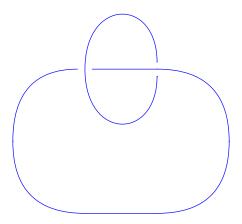
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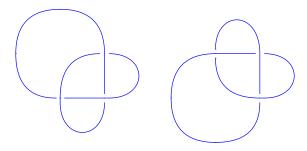
(continued from previous page)

Graphics object consisting of ... graphics primitives



Plotting links with multiple isolated components:

```
sage: L = Link([[[-1, 2, -3, 1, -2, 3], [4, -5, 6, -4, 5, -6]], [1, 1, 1, 1, 1, 1, 1, 1])
sage: L.plot()
Graphics object consisting of ... graphics primitives
```



If a coloring is passed, the different arcs are plotted with the corresponding colors:

```
sage: B = BraidGroup(4)
sage: b = B([1,2,3,1,2,-1,-3,2,3])
sage: L = Link(b)
sage: L.plot(color=L.colorings(3)[0])
Graphics object consisting of ... graphics primitives
```

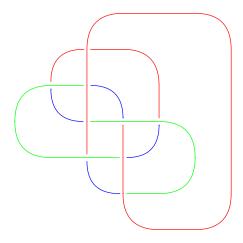
regions()

Return the regions from the link diagram of self.

Regions are obtained always turning left at each crossing.

Then the regions are represented as a list with the segments that form its boundary, with a sign depending on the orientation of the segment as part of the boundary.

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EXAMPLES:

```
\rightarrow -1, -1, 1, -1, 1]])
sage: L.regions()
[[14, -5, 12, -9], [13, 9], [11, 5, 1, 7, 3], [10, -3, 8, -13], [6, -1], [4, -1], [10, -3, 10, -3]
\hookrightarrow11], [2, -7], [-2, -6, -14, -8], [-4, -10, -12]]
sage: L = Link([[[1, -2, 3, -4, 2, -1, 4, -3]],[1, 1, -1, -1]])
sage: L.regions()
[[8, 4], [7, -4, 1], [6, -1, -3], [5, 3, -8], [2, -5, -7], [-2, -6]]
sage: L = Link([[-1, +2, 3, -4, 5, -6, 7, 8, -2, -5, +6, +1, -8, -3, 4, -7]],[-
\rightarrow 1, -1, -1, -1, 1, -1, 1]])
sage: L.regions()
[[16, 8, 14, 4], [15, -4], [13, -8, 1], [12, -1, -7], [11, 7, -16, 5], [10, -5]
\hookrightarrow -15, -3], [9, 3, -14], [6, -11], [2, -9, -13], [-2, -12, -6, -10]]
sage: B = BraidGroup(2)
sage: L = Link(B([-1, -1, -1]))
sage: L.regions()
[[6, -5], [5, 1, 3], [4, -3], [2, -1], [-2, -6, -4]]
sage: L = Link([[[1, -2, 3, -4], [-1, 5, -3, 2, -5, 4]], [-1, 1, 1, -1, -1]])
sage: L.regions()
[[10, -4, -7], [9, 7, -3], [8, 3], [6, -9, -2], [5, 2, -8, 4], [1, -5], [-1, -9]
\rightarrow10, -6]]
sage: L = Link([[1, 2, 3, 3], [2, 5, 4, 4], [5, 7, 6, 6], [7, 1, 8, 8]])
sage: L.regions()
[[-3], [-4], [-6], [-8], [7, 1, 2, 5], [-1, 8, -7, 6, -5, 4, -2, 3]]
```

Note: The link diagram is assumed to have only one completely isolated component. This is because otherwise some regions would have disconnected boundary.

seifert_circles()

Return the Seifert circles from the link diagram of self.

Seifert circles are the circles obtained by smoothing all crossings respecting the orientation of the segments.

Each Seifert circle is represented as a list of the segments that form it.

```
sage: L = Link([[[1, -2, 3, -4, 2, -1, 4, -3]], [1, 1, -1, -1]])
sage: L.seifert_circles()
[[1, 7, 5, 3], [2, 6], [4, 8]]
sage: L = Link([[-1, 2, 3, -4, 5, -6, 7, 8, -2, -5, 6, 1, -8, -3, 4, -7]], [-1, -1, -2, -3, -4, -7]]
\rightarrow -1, -1, -1, 1, 1, -1, 1]])
sage: L.seifert_circles()
[[1, 13, 9, 3, 15, 5, 11, 7], [2, 10, 6, 12], [4, 16, 8, 14]]
sage: L = Link([[[-1, 2, -3, 4, 5, 1, -2, 6, 7, 3, -4, -7, -6, -5]], [-1, -1, -
\hookrightarrow 1, -1, 1, -1, 1]])
sage: L.seifert_circles()
[[1, 7, 3, 11, 5], [2, 8, 14, 6], [4, 12, 10], [9, 13]]
sage: L = Link([[1, 7, 2, 6], [7, 3, 8, 2], [3, 11, 4, 10], [11, 5, 12, 4], [14,
\rightarrow 5, 1, 6], [13, 9, 14, 8], [12, 9, 13, 10]])
sage: L.seifert_circles()
[[1, 7, 3, 11, 5], [2, 8, 14, 6], [4, 12, 10], [9, 13]]
sage: L = Link([[[-1, 2, -3, 5], [4, -2, 6, -5], [-4, 1, -6, 3]], [-1, 1, 1, 1, 1, 1]
\hookrightarrow -1, -1]])
sage: L.seifert_circles()
[[1, 11, 8], [2, 7, 12, 4, 5, 10], [3, 9, 6]]
sage: B = BraidGroup(2)
sage: L = Link(B([1, 1, 1]))
sage: L.seifert_circles()
[[1, 3, 5], [2, 4, 6]]
```

seifert_matrix()

Return the Seifert matrix associated with self.

ALGORITHM:

This is the algorithm presented in Section 3.3 of [Col2013].

OUTPUT

The intersection matrix of a (not necessarily minimal) Seifert surface.

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, 3, 1, 3]))
sage: L.seifert_matrix()
[ 0   0]
[ 0 -1]
sage: B = BraidGroup(8)
sage: L = Link(B([-1, 3, 1, 5, 1, 7, 1, 6]))
sage: L.seifert_matrix()
[ 0   0   0]
[ 1 -1   0]
[ 0   1 -1]
sage: L = Link(B([-2, 4, 1, 6, 1, 4]))
sage: L.seifert_matrix()
[-1   0]
[ 0 -1]
```

signature()

Return the signature of self.

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This is defined as the signature of the symmetric matrix

$$V + V^t$$
,

where V is the Seifert matrix.

See also:

```
omega_signature(), seifert_matrix()
```

EXAMPLES:

```
sage: B = BraidGroup(4)
sage: L = Link(B([-1, 3, 1, 3]))
sage: L.signature()
-1
sage: B = BraidGroup(8)
sage: L = Link(B([-2, 4, 1, 6, 1, 4]))
sage: L.signature()
-2
sage: L = Link(B([1, 2, 1, 2]))
sage: L.signature()
-2
```

writhe()

Return the writhe of self.

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CHAPTER

THREE

ACCESS TO THE KNOTINFO DATABASE

This module contains the class *KnotInfoBase* which is derived from Enum and provides knots and links listed in the databases at the web-pages KnotInfo and LinkInfo as its items.

This interface contains a set of about twenty knots and links statically as demonstration cases. The complete database can be installed as an optional Sage package using

- sage -i database_knotinfo (does not install if the current version is present)
- sage -f database_knotinfo (installs even if the current version is present)

To perform all the doctests concerning the usage of the database on the installation add the option -c. In this case (for instance sage -f -c database_knotinfo) the installation breaks on failing tests.

The installation of the complete database will be necessary in order to have access to all the properties recorded in the databases, as well.

If the entire database is installed as explained above, the import instructions for *KnotInfo* and *KnotInfoSeries*, which can be seen in the opening lines of the examples, are unnecessary.

Be aware that there are a couple of conventions used differently on KnotInfo as in Sage, especially concerning the selection of the symmetry version of the link.

In this context you should note that the PD notation is recorded counter clockwise in KnotInfo (see note in *KnotInfoBase.link()*). In our transition to Sage objects this is translated (by default) in order to avoid confusion about exchanged mirror versions.

Also, note that the braid notation is used according to Sage, even thought in the source where it is taken from, the braid generators are assumed to have a negative crossing which would be opposite to the convention in Sage (see definition 3 of Gittings, T., "Minimum Braids: A Complete Invariant of Knots and Links).

For different conventions regarding normalization of the polynomial invariants see the according documentation of KnotInfoBase.homfly_polynomial(), KnotInfoBase.jones_polynomial() and KnotInfoBase.alexander_polynomial().

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.pd_notation()
[[6, 1, 7, 2], [8, 3, 5, 4], [2, 5, 3, 6], [4, 7, 1, 8]]
sage: L.pd_notation(original=True)
'{{6, 1, 7, 2}, {8, 3, 5, 4}, {2, 5, 3, 6}, {4, 7, 1, 8}}'
sage: L.is_knot()
False
sage: L.num_components()
2
```

Items for knots need a leading K for technical reason:

```
sage: K = KnotInfo.K4_1
sage: K.is_knot()
True
```

Injecting the variable name into the namespace:

```
sage: KnotInfo.K5_1.inject()
Defining K5_1
sage: K5_1.dt_notation()
[6, 8, 10, 2, 4]
```

Defining a link from the original name string:

```
sage: KnotInfo('L6a1{1}').inject()
Defining L6a1_1
sage: L6a1_1.is_alternating()
True
```

Obtaining an instance of Braid:

```
sage: L.braid()
s0*s1^-1*s2*s1^-1*s0^-1*s1^-1*s2^-1*s1^-1
sage: type(_)
<class 'sage.groups.braid.BraidGroup_class_with_category.element_class'>
```

Obtaining an instance of *Link*:

```
sage: 1 = L.link(); 1
Link with 2 components represented by 4 crossings
sage: type(1)
<class 'sage.knots.link.Link'>
```

If you have SnapPy installed inside Sage you can obtain an instance of Link, too:

```
sage: L6 = KnotInfo.L6a1_0
sage: 16s = L6.link(snappy=True); 16s
                                          # optional - snappy
Plink failed to import tkinter.
<Link: 2 comp; 6 cross>
sage: type(16s)
                                           # optional - snappy
<class 'spherogram.links.invariants.Link'>
sage: 16 = L6.link().mirror_image()
sage: 16 == 16s.sage_link()
                                           # optional - snappy
True
sage: L6.link(L6.items.name, snappy=True) # optional - snappy
<Link L6a1: 2 comp; 6 cross>
sage: 16sn = _
                                           # optional - snappy
sage: 16s == 16sn
                                           # optional - snappy
False
sage: l6sn.sage_link().is_isotopic(16)
                                           # optional - snappy
```

But observe that the name conversion to SnapPy does not distingiush orientation types:

```
sage: L6b = KnotInfo.L6a1_1
sage: L6b.link(L6b.items.name, snappy=True) # optional - snappy
<Link L6a1: 2 comp; 6 cross>
sage: _.PD_code() == l6sn.PD_code() # optional - snappy
True
```

Obtaining the HOMFLY-PT polynomial:

```
sage: L.homfly_polynomial()
-v^-1*z - v^-3*z - v^-3*z^-1 + v^-5*z^-1
sage: _ == l.homfly_polynomial(normalization='vz')
True
```

Obtaining the original string from the database for an arbitrary property:

```
sage: K[K.items.classical_conway_name] # optional - database_knotinfo
'4_1'
```

Further methods:

```
sage: K.crossing_number()
sage: K.gauss_notation()
[-1, 2, -3, 1, -4, 3, -2, 4]
sage: K.dt_notation()
[4, 6, 8, 2]
sage: K.determinant()
sage: K.symmetry_type()
'fully amphicheiral'
sage: _ == K[K.items.symmetry_type]
True
sage: K.is_reversible()
sage: K.is_amphicheiral()
sage: K.jones_polynomial()
t^2 - t - 1/t + 1/t^2 + 1
sage: K.kauffman_polynomial()
a^2z^2 + a^2 - a^2 - a^2 - a^2 + 2^2 + a^{-1}z^3 - 1 - a^{-1}z + a^{-2}z^2 - a^{-2}z^2
sage: K.alexander_polynomial()
t^2 - 3*t + 1
```

Using the column_type of a property:

You can launch web-pages attached to the links:

```
sage: K.diagram()  # not tested
True
sage: L.diagram(single=True)  # not tested
True
sage: L.knot_atlas_webpage()  # not tested
True
sage: K.knotilus_webpage()  # not tested
True
```

and the description web-pages of the properties:

```
sage: K.items.positive.description_webpage() # not tested
True
```

To see all the properties available in this interface you can use "tab-completion". For example type K.items. and than hit the "tab-key". You can select the item you want from the list. If you know some first letters type them first to obtain a reduced selection list.

In a similar way you may select the knots and links. Here you have to type KnotInfo. or KnotInfo.L7 before stroking the "tab-key". In the latter case the selection list will be reduced to proper links with 7 crossings.

Finally there is a method Link. get_knotinfo() of class Link to find an instance in the KnotInfo database:

REFERENCES:

- KnotInfo
- LinkInfo

AUTHORS:

Sebastian Oehms August 2020: initial version

Thanks to Chuck Livingston and Allison Moore for their support. For further acknowledgments see the correspondig hompages.

class sage.knots.knotinfo.KnotInfo

```
Bases: sage.knots.knotinfo.KnotInfoBase
```

An enumeration.

class sage.knots.knotinfo.KnotInfoBase

```
Bases: enum. Enum
```

Enum class to select the knots and links listed in the databases at the web-pages KnotInfo and LinkInfo.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: [knot.name for knot in KnotInfo if knot.crossing_number() < 5]
['K0_1', 'K3_1', 'K4_1', 'L2a1_0', 'L2a1_1', 'L4a1_0', 'L4a1_1']</pre>
```

More examples and information can be seen in the module header *knotinfo* (by typing):

```
sage: import sage.knots.knotinfo # not tested
sage: sage.knots.knotinfo? # not tested
```

alexander_polynomial(var='t', original=False, laurent_poly=False)

Return the Alexander polynomial according to the value of column alexander_polynomial for this knot as an instance of Polynomial.

It is obtained from the Seifert matrix V of self by the following formula (see the KnotInfo description web-page; to launch it see the example below):

$$A(L) = \det(V - tV^t)$$

Here V^t stands for the transpose of V.

INPUT:

- var (default: 't') the variable
- original boolean (optional, default False) if set to True the original table entry is returned as a string
- laurent_poly boolean (default False) see the note below

OUTPUT:

A polynomial over the integers, more precisely an instance of Polynomial. If laurent_poly is set to True a Laurent polynomial over the integers, more precisely an instance of LaurentPolynomial is returned. If original is set to True then a string is returned.

Note: As an invariant the Alexander polynomial is only unique up to a unit factor in the Laurent polynomial ring over the integers in the indeterminate t. While the normalization of the exponents in KnotInfo guarantees it to be a proper polynomial, this is not the case for the implementation in Sage. Use the keyword laurent_poly to achiev a normalization according to Sage. But, still there may be a difference in sign (see the example below).

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K4_1
sage: Ka = K.alexander_polynomial(); Ka
t^2 - 3*t + 1
```

Comparison with Sage's results:

```
sage: k = K.link()
sage: ka = k.alexander_polynomial(); ka
-t^-1 + 3 - t
sage: K.alexander_polynomial(laurent_poly=True)
t^-1 - 3 + t
sage: _ == -ka
True
```

Launch the KnotInfo description web-page:

```
sage: K.items.alexander_polynomial.description_webpage() # not tested
True
```

braid()

Return the braid notation of self as an instance of Braid.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.braid()
s^3
sage: K.braid_notation()
(1, 1, 1)
```

braid_index()

Return the value of column braid_index for this link as a Python int.

OUTPUT:

Python int giving the minimum of strands needed to represent self as closure of a braid.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.braid_index()
4
```

braid_length()

Return the value of column braid_length for this link as a Python int.

OUTPUT:

Python int giving the minimum length of a braid word needed to represent self as closure of a braid.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.braid_length()
3
```

braid_notation(original=False)

Return the value of column braid_notation for this link as a Python tuple (Tietze form). For more information type KnotInfo.K0_1.items.braid_notation.description_webpage().

INPUT:

• original – boolean (optional, default False) if set to True the original table entry is returned as a string

OUTPUT:

Python tuple representing the braid whose closure is self in Tietze form.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.braid_notation()
(1, -2, 3, -2, -1, -2, -3, -2)
```

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```
sage: L.braid_notation(original=True)
'{4, {1, -2, 3, -2, -1, -2, -3, -2}}'
```

crossing_number()

Return the minimal number of crossings of self.

Note: In contrast to the number of crossings displayed for instances of *Link* this number is the minimum over all possible diagrams of the link. The number of crossings displayed in the representation string of *Link* refers to the special diagram which could be larger.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L4a1_0.crossing_number()
4
sage: KnotInfo.K3_1.crossing_number()
3
sage: Link(KnotInfo.L4a1_0.braid())
Link with 2 components represented by 8 crossings
```

determinant()

Return the determinant of self.

From the KnotInfo description page:

The determinant of a knot is $det(V + V^t)$, where V is a Seifert matrix for the knot.

To read the complete description type KnotInfo.K0_1.items.determinant.description_webpage().

Note: KnotInfo's value for the unknot **0**_1 is zero. This is not compatible whith Sage's result (the value of the Alexander polynomial at -1). Since this method is needed to identify Sage links we take the according value in that case.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L4a1_0.determinant()
4
sage: KnotInfo.K3_1.determinant()
3
sage: KnotInfo.K0_1.determinant()
1
```

diagram(single=False, new=0, autoraise=True)

Launch the diagram of self given on the KnotInfo web-page.

INPUT:

- single boolean (default False) if set to True only one diagram is shown.
- new int according to open() of webbrowser (0 default, 1 new window, 2 new tab)
- autoraise boolean (default True)

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.diagram()  # not tested
True
sage: K.diagram(single=True) # not tested
True
```

dt_notation(original=False)

Return the value of column dt_notation for this link as a Python list of Python lists. For more information type KnotInfo.K0_1.items.dt_notation.description_webpage().

INPUT:

• original – boolean (optional, default False) if set to True the original table entry is returned as a string

OUTPUT:

Python list of python lists each entry of the outer list representing a crossing.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.dt_notation()
[[6, 8], [2, 4]]
sage: L.dt_notation(original=True)
'[{6, 8}, {2, 4}]'
sage: L = KnotInfo.L4a1_0
sage: K = KnotInfo.K4_1
sage: K.dt_notation()
[4, 6, 8, 2]
```

gauss_notation(original=False)

Return the value of column gauss_notation for this link as a Python list of Python lists. For more information type KnotInfo.K0_1.items.gauss_notation.description_webpage().

INPUT:

• original – boolean (optional, default False) if set to True the original table entry is returned as a string

OUTPUT:

Python list of

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.gauss_notation()
[[1, -3, 2, -4], [3, -1, 4, -2]]
sage: L.gauss_notation(original=True)
'{{1, -3, 2, -4}, {3, -1, 4, -2}}'
```

homfly_polynomial(var1='v', var2='z', original=False)

Return the HOMFLY-PT polynomial according to the value of column homfly_polynomial for this knot or link (in the latter case the column homflypt_polynomial is used) as an instance of the element class

according to the output of Link.homfly_polynomial() of Link.

The HOMFLY-PT polynomial P(L) of a link L satisfies the following skein relation (see the corresponding KnotInfo description page):

$$P(O) = 1, \ v^{-1}P(L_{+}) - vP(L_{-}) = zP(L_{0})$$

INPUT:

- var1 string (default v) for the name of the first variable
- var2 string (default z) for the name of the second variable
- original boolean (default False) if set to True the original table entry is returned as a string

OUTPUT:

A Laurent polynomial over the integers, more precisely an instance of LaurentPolynomial_mpair. If original is set to True then a string is returned.

Note: The skein-relation for the HOMFLY-PT polynomial given on KnotInfo does not match the default used in Sage. For comparison you have to use the keyword argument normalization='vz' on the side of Sage.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K3_1 = KnotInfo.K3_1
sage: PK3_1 = K3_1.homfly_polynomial(); PK3_1
-v^4 + v^2*z^2 + 2*v^2
sage: K3_1.homfly_polynomial(original=True)
'(2*v^2-v^4)+(v^2)*z^2'
sage: PK3_1 == K3_1.link().homfly_polynomial(normalization='vz')
True
```

for proper links:

```
sage: L4a1_1 = KnotInfo.L4a1_1
sage: PL4a1_1 = L4a1_1.homfly_polynomial(var1='x', var2='y'); PL4a1_1
-x^5*y + x^3*y^3 - x^5*y^-1 + 3*x^3*y + x^3*y^-1
sage: _ == L4a1_1.link().homfly_polynomial('x', 'y', 'vz')
True
```

check the skein-relation from the KnotInfo description page (applied to one of the positive crossings of the right-handed trefoil):

```
sage: R = PK3_1.parent()
sage: P0 = R.one()
sage: L2a1_1 = KnotInfo.L2a1_1
sage: PL2a1_1 = L2a1_1.homfly_polynomial()
sage: v, z = R.gens()
sage: ~v*PK3_1 -v*P0 == z*PL2a1_1
True
```

REFERENCES:

• Wikipedia article HOMFLY_polynomial

inject(verbose=True)

Inject self with its name into the namespace of the Python code from which this function is called.

INPUT:

• verbose – boolean (optional, default True) to suppress the message printed on the invocation

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.inject()
Defining K5_2
sage: K5_2.is_alternating()
True
```

is_adequate()

Return whether self is adequate.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_adequate() # optional - database_knotinfo
True
```

is_almost_alternating()

Return whether self is almost alternating.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_almost_alternating() # optional - database_knotinfo
False
```

is_alternating()

Return whether self is alternating.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_alternating()
True
```

is_amphicheiral(positive=False)

Return whether self is amphicheiral.

INPUT:

• positive – boolean (optional, default False) whether to check if self is positive or negative amphicheiral (see documentation of symmetry_type())

OUTPUT:

Boolean or None if this cannot be determined.

True if self is fully or negative amphicheiral per default. If positive is set to True than fully and positive amphicheiral links give True.

Note: For proper links this property is not provided in the database. Anyway, we support it here in this case, as well, except for a few items where it cannot be determined easily and where None is returned as

answer.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: Kp = KnotInfo.K12a_427
                                         # optional - database_knotinfo
sage: Kp.is_amphicheiral()
                                         # optional - database_knotinfo
False
sage: Kp.is_amphicheiral(positive=True) # optional - database_knotinfo
True
sage: Kn = KnotInfo.K10_88
                                         # optional - database_knotinfo
sage: Kn.is_amphicheiral()
                                         # optional - database_knotinfo
True
sage: Kn.is_amphicheiral(positive=True) # optional - database_knotinfo
False
sage: KnotInfo.L4a1_0.is_amphicheiral()
False
sage: KnotInfo.L10n59_1.is_amphicheiral() # optional - database_knotinfo
True
sage: KnotInfo.L10n59_0.inject()
                                          # optional - database_knotinfo
Defining L10n59_0
sage: L10n59_0.is_amphicheiral() is None # optional - database_knotinfo
True
```

is_fibered()

Return whether self is fibered.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K6_3.is_fibered()
True
```

is_knot()

Return whether self is a knot or a proper link.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L7a1_0.is_knot() # optional - database_knotinfo
False
sage: KnotInfo.K6_3.is_knot()
True
```

is_oriented()

Return whether self is oriented.

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L6a2_1.is_oriented()
True
```

is_positive()

Return whether self is positive.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_positive()
True
```

is_positive_braid()

Return whether self is a positive braid.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_positive_braid() # optional - database_knotinfo
False
```

is_quasi_alternating()

Return whether self is quasi alternating.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_quasi_alternating() # optional - database_knotinfo
True
```

is_quasipositive()

Return whether self is quasi-positive.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_quasipositive() # optional - database_knotinfo
True
```

is_recoverable(unique=True)

Return if self can be recovered from its conversion to Sage links using the pd_notation and the braid_notation and their mirror images.

The method is indirectly used by the TestSuite of the series of self.

INPUT:

• unique – boolean (optional, default=``True``) if set to False it is only checked if self is among the recovered items

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L4a1_0.inject()
Defining L4a1_0
sage: L4a1_0.is_recoverable()
True
sage: L4a1_0.is_recoverable(unique=False)
True
sage: KnotInfo.L5a1_0.inject()
Defining L5a1_0
```

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```
sage: L5a1_0.is_recoverable()
False
sage: L5a1_0.is_recoverable(unique=False)
True
```

is_reversible()

Return whether self is reversible.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K6_3.is_reversible()
True
```

is_strongly_quasipositive()

Return whether self is strongly quasi-positive.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.is_strongly_quasipositive() # optional - database_knotinfo
True
```

is_unique()

Return whether there is no other isotopic link in the database or not.

OUTPUT:

Boolean or None if this cannot be determined.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L4a1_0.is_unique()
True
sage: KnotInfo.L5a1_0.is_unique()
False
sage: L = KnotInfo.L7a7_0_0  # optional - database_knotinfo
sage: L.series(oriented=True).inject()  # optional - database_knotinfo
Defining L7a7
sage: [(L,L.is_unique()) for L in L7a7]  # optional - database_knotinfo
[(<KnotInfo.L7a7_0_0: 'L7a7{0,0}'>, False),
  (<KnotInfo.L7a7_1_0: 'L7a7{1,0}'>, None),
  (<KnotInfo.L7a7_0_1: 'L7a7{0,1}'>, False),
  (<KnotInfo.L7a7_0_1: 'L7a7{0,1}'>, True)]
```

items

Return an Enum class to select a column item of the KnotInfo database.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: it = L.items
sage: [i.name for i in it if i.name.startswith('braid')]
['braid_index', 'braid_length', 'braid_notation']
```

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```
sage: L.items.dt_notation.column_name()
'DT Notation'
```

To check if the item is available for proper links or only knots type:

```
sage: it.gauss_notation.column_type()
<KnotInfoColumnTypes.KnotsAndLinks: 'B'>
sage: it.dt_notation.column_type()
<KnotInfoColumnTypes.OnlyKnots: 'K'>
```

To see the description of the item in your web browser type:

```
sage: it.gauss_notation.description_webpage() # not tested
True
```

Return the Jones polynomial according to the value of column jones_polynomial for this knot or link as an element of the symbolic ring SR or an instance of LaurentPolynomial depending on the keyword skein_normalization. Using the keyword puiseux instead of an element of the symbolic ring an instance of PuiseuxSeries can be returned.

The Jones polynomial V(L) of a link L satisfies the following skein relation (see the corresponding Knot-Info description page):

$$V(O) = 1, \ t^{-1}V(L_{+}) - tV(L_{-}) = (t^{\frac{1}{2}} - t^{-\frac{1}{2}})V(L_{0})$$

INPUT:

- variab variable (default: None) used according to Link.jones_polynomial()
- skein_normalization boolean (default: False) used according to Link.jones_polynomial()
- puiseux boolean (default True) only used in case skein_normalization=False. If set to True instead of an element of the symbolic ring an instance of PuiseuxSeries is returned
- original boolean (default False) if set to True the original table entry is returned as a string
- use_sqrt boolean (default False) see the note below

OUTPUT:

Depends on the keywords (in excluding order):

- original=True a string according to the original value from the database
- skein_normalization=True a Laurent polynomial over the integers, more precisely an instance of LaurentPolynomial
- puiseux=True a puiseux series over the integers, more precisely an instance of PuiseuxSeries

In all other cases an element of the symbolic ring SR.

Note: There is a difference to Sage's conventions concerning the Jones polynomial in the case of proper links. KnotInfo does not display these polynomials in the indeterminate t used in the skein relation. Instead a variable x is used defined by $x^2 = t$. Sage uses t in both cases, knots and proper links. Thus, to obtain the Jones polynomial for a proper link in t you have to set the keyword use_sqrt to True.

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K4_1
sage: Kj = K.jones_polynomial(); Kj
t^2 - t - 1/t + 1/t^2 + 1
sage: Kjs = K.jones_polynomial(skein_normalization=True); Kjs
A^-8 - A^-4 + 1 - A^4 + A^8
sage: Kjp = K.jones_polynomial(puiseux=True); Kjp
t^-2 - t^-1 + 1 - t + t^2
```

for proper links:

```
sage: L = KnotInfo.L2a1_1
sage: Lj = L.jones_polynomial(); Lj
-x^5 - x
sage: Ljt = L.jones_polynomial(use_sqrt=True); Ljt
-t^{(5/2)} - sqrt(t)
sage: Ljp = L.jones_polynomial(puiseux=True); Ljp
-t^{(1/2)} - t^{(5/2)}
sage: Ljs = L.jones_polynomial(skein_normalization=True); Ljs
-A^2 - A^10
sage: Lj.parent()
Symbolic Ring
sage: Ljt.parent()
Symbolic Ring
sage: Ljp.parent()
Puiseux Series Ring in t over Integer Ring
sage: Ljs.parent()
Univariate Laurent Polynomial Ring in A over Integer Ring
```

Comparison with Sage's results:

```
sage: k = K.link()
sage: kj = k.jones_polynomial()
sage: bool(Kj == kj)
True
sage: kjs = k.jones_polynomial(skein_normalization=True)
sage: Kjs == kjs
True
sage: l = L.link()
sage: lj = 1.jones_polynomial()
sage: bool(Lj == lj)
False
sage: bool(Ljt == lj)
                        # see note above
sage: ljs = 1.jones_polynomial(skein_normalization=True)
sage: Ljs == ljs
True
```

Check the skein-relation from the KnotInfo description page (applied to one of the positive crossings of the right-handed trefoil):

```
sage: K3_1 = KnotInfo.K3_1
sage: K3_1j = K3_1.jones_polynomial()
```

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```
sage: L2a1_1j = Ljt  # see note above
sage: R = L2a1_1j.parent()
sage: Oj = R(1)
sage: t = R('t')
sage: lhs = expand(~t*K3_1j - t*Oj)
sage: rhs = expand((sqrt(t) - ~sqrt(t))*L2a1_1j)
sage: bool(lhs == rhs)
True
```

The same with the Puiseux series version:

```
sage: K3_1jp = K3_1.jones_polynomial(puiseux=True)
sage: L2a1_1jp = Ljp
sage: R = L2a1_1jp.parent()
sage: Ojp = R(1)
sage: t = R('t')
sage: ~t*K3_1jp - t*Ojp == (t^(1/2)-~t^(1/2))*L2a1_1jp
True
```

The same in the case of skein normalization (using $t = A^4$):

```
sage: K3_1js = K3_1.jones_polynomial(skein_normalization=True)
sage: L2a1_1js = L.jones_polynomial(skein_normalization=True)
sage: Rs = K3_1js.parent()
sage: Ojs = Rs.one()
sage: A, = Rs.gens()
sage: ~A^4*K3_1js - A^4*Ojs == (A^2-~A^2)*L2a1_1js
True
```

REFERENCES:

• Wikipedia article Jones_polynomial

kauffman_polynomial(*var1='a'*, *var2='z'*, *original=False*)

Return the Kauffman polynomial according to the value of column kauffman_polynomial for this knot or link as an instance of LaurentPolynomial_mpair.

The Kauffman polynomial F(L) respectively its corresponding invariant under regular isotopy $\Delta(L) = a^{w(L)}F(L)$ where w(L) is the writhe of the link L satisfies the following skein relation (see the corresponding KnotInfo description page):

$$\Delta(O) = 1$$
, $\Delta(L_+) - \Delta(L_-) = z(\Delta(L_0 + \Delta(L_\infty)))$

Furthermore, removing a curl of sign ϵ leads to a multiplication of $\Delta(L)$ with a^{ϵ} .

INPUT:

- var1 (default: 'a') the first variable
- var2 (default: 'z') the second variable
- original boolean (optional, default False) if set to True the original table entry is returned as a string

OUTPUT:

A Laurent polynomial over the integers, more precisely an instance of LaurentPolynomial_mpair. If original is set to False then a string is returned.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L2a1_1
sage: L.kauffman_polynomial()
a^-1*z - a^-1*z^-1 + a^-2 + a^-3*z - a^-3*z^-1
sage: K = KnotInfo.K4_1
sage: K.kauffman_polynomial()
a^2*z^2 + a*z^3 - a^2 - a*z + 2*z^2 + a^-1*z^3 - 1 - a^-1*z + a^-2*z^2 - a^-2
```

Comparison with Jones polynomial:

```
sage: k = _
sage: a, z = k.variables()
sage: j = K.jones_polynomial(skein_normalization=True)
sage: t, = j.variables()
sage: k.subs(a=-t^3, z=~t+t) == j.subs(t=t^4)
True
```

Check the skein relation:

```
sage: K3_1 = KnotInfo.K3_1
sage: FK3_1 = K3_1.kauffman_polynomial()
sage: FL2a1_1 = L.kauffman_polynomial()
sage: z, a = FK3_1.variables()
sage: \triangleK3_1 = FK3_1 * a**K3_1.link().writhe()
sage: \triangleL2a1_1 = FL2a1_1 * a**L.link().writhe()
sage: \triangleO1p = a # unknot with one positive curl
sage: \triangleO2n = a**-2 # unknot with two negative curls
sage: \triangleK3_1 + \triangleO1p == z*(\triangleL2a1_1 + \triangleO2n)
True
```

REFERENCES:

• Wikipedia article Kauffman_polynomial

knot_atlas_webpage(new=0, autoraise=True)

Launch the Knot Atlas web-page for self.

INPUT:

- new int according to open() of webbrowser (0 default, 1 new window, 2 new tab)
- autoraise boolean (default True)

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.knot_atlas_webpage() # not tested
True
```

knotilus_webpage(new=0, autoraise=True)

Launch the Knotilus web-page for self.

INPUT:

• new – int according to open() of webbrowser (0 default, 1 new window, 2 new tab)

• autoraise – boolean (default True)

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.knotilus_webpage(new=1) # not tested
True
```

link(use_item=<Columns.pd_notation: ['PD Notation', <KnotInfoColumnTypes.OnlyKnots: 'K'>]>, snappy=False)

Return self as an instance of Link or optional spherogram.links.invariants.Link (SnapPy).

INPUT:

- use_item (optional, default self.items.pd_notation) instance of KnotInfoColumns to choose the column that should be used to construct the link. Allowed values are:
 - self.items.pd_notation
 - self.items.braid_notation
 - self.items.name (only for snappy=True)
 - self.items.dt_notation (only for knots and snappy=False)
 - self.items.gauss_notation (only for knots and snappy=False)
- snappy boolean (default False) if set to True the target of the conversion is the pip installable package SnapPy (explicitly, spherogram.links.invariants.Link). If SnapPy is not installed an ImportError is raised. To install SnapPy use sage -pip install snappy.

Note: We use the PD-notation to construct self as default. This ensures that the number of crossings displayed in the representation string of the link coincides with the crossing number as a topological invariant.

But attention: The convention on how the edges are listed are opposite to each other

- KnotInfo: counter clockwise
- · Sage: clockwise

Therefore, we take the mirror version of the pd_notation!

Furthermore, note that the mirror version may depend on the used KnotInfo-notation. For instance, regarding to the knot 5_1 the Gauss- and the DT-notation refer to the mirror image (see example below).

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K = KnotInfo.K3_1
sage: K.link()
Knot represented by 3 crossings
sage: _.braid()
s^3
sage: _ == K.braid()
True
```

using dt_notation:

```
sage: K.link(use_item=K.items.dt_notation)
Knot represented by 3 crossings
sage: _.braid()
s^3

sage: L = KnotInfo.L4a1_0
sage: L.link()
Link with 2 components represented by 4 crossings

sage: L.link(use_item=L.items.dt_notation)
Traceback (most recent call last):
...
ValueError: Link construction using Columns.dt_notation not possible
```

using snappy:

but observe:

using braid_notation:

```
sage: L2.link(use_item=L.items.braid_notation) == 12
True
```

observe:

```
sage: L.link(use_item=L.items.braid_notation)
Link with 2 components represented by 8 crossings

sage: K6_1 = KnotInfo.K6_1
sage: K6_1.link().braid() == K6_1.braid()
False
```

also observe:

```
sage: K4_1 = KnotInfo.K4_1
sage: K4_1.link().pd_code()
[[4, 1, 5, 2], [8, 5, 1, 6], [6, 4, 7, 3], [2, 8, 3, 7]]
sage: K4_1.pd_notation()
[[4, 2, 5, 1], [8, 6, 1, 5], [6, 3, 7, 4], [2, 7, 3, 8]]
sage: K5_1 = KnotInfo.K5_1
sage: K5_1.link().braid()
s^5
sage: K5_1.link(K5_1.items.dt_notation).braid()
s^-5
sage: K5_1.link(K5_1.items.gauss_notation).braid()
s^-5
```

name_unoriented()

Return the the part of the name of self which is independent on the orientation.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L10a122_1_0.name_unoriented() # optional - database_knotinfo
'L10a122'
```

num_components()

Return the number of components of self.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L6a1_0.num_components()
2
```

pd_notation(original=False)

Return the value of column pd_notation for this link as a Python list of Python lists. For more information type KnotInfo.K0_1.items.pd_notation.description_webpage().

INPUT:

• original – boolean (optional, default False) if set to True the original table entry is returned as a string

OUTPUT:

Python list of python lists each entry of the outer list representing a crossing.

```
sage: from sage.knots.knotinfo import KnotInfo
sage: L = KnotInfo.L4a1_0
sage: L.pd_notation()
[[6, 1, 7, 2], [8, 3, 5, 4], [2, 5, 3, 6], [4, 7, 1, 8]]
sage: L.pd_notation(original=True)
'{{6, 1, 7, 2}, {8, 3, 5, 4}, {2, 5, 3, 6}, {4, 7, 1, 8}}'
sage: K = KnotInfo.K4_1
sage: K.pd_notation()
[[4, 2, 5, 1], [8, 6, 1, 5], [6, 3, 7, 4], [2, 7, 3, 8]]
```

series(oriented=False)

Return the series of links self belongs to.

INPUT:

• oriented – boolean (default False) it only affects proper links. By default the items of the series will be again series of links collecting all orientation mutants of an unoriented name. To obtain the series of the individual links this keyword has to be set to True.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: K5 = KnotInfo.K5_2.series()
sage: K5(1)
<KnotInfo.K5_1: '5_1'>
sage: KnotInfo.L4a1_1.series().inject()
Defining L4a
sage: L4a(1)
Series of links L4a1
sage: KnotInfo.L4a1_1.series(oriented=True).inject()
Defining L4a1
sage: L4a(1) == L4a1
True
sage: L4a1(1)
<KnotInfo.L4a1_1: 'L4a1{1}'>
```

signature()

Return the signature of self.

From the KnotInfo description page:

The signature of a knot, $\sigma(K)$, is equal to $\sigma(V+V^t)$, the signature of $V+V^t$ where V is a Seifert matrix for the knot and V^t is its transpose.

To read the complete description type KnotInfo.K0_1.items.signatur.description_webpage().

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.signature() # optional - databsase_knotinfo
1
```

symmetry_type()

Return the symmetry type of self.

From the KnotInfo description page:

If a knot is viewed as the oriented diffeomorphism class of an oriented pair, $K=(S_3,S_1)$, with S_i diffeomorphic to S^i , there are four oriented knots associated to any particular knot K. In addition to K itself, there is the reverse, $K^r=(S_3,-S_1)$, the concordance inverse, $-K=(-S_3,-S_1)$, and the mirror image, $K^m=(-S_3,S_1)$. A knot is called reversible if $K=K^r$, negative amphicheiral if $K=K^r$, and positive amphicheiral if $K=K^r$.

A knot possessing any two of these types of symmetry has all three. Thus, in the table, a knot is called reversible if that is the only type of symmetry it has, and likewise for negative amphicheiral. If it has none of these types of symmetry it is called chiral, and if it has all three it is called fully amphicheiral.

For prime knots with fewer than 12 crossings, all amphicheiral knots are negative amphicheiral.

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K6_1.series().inject()
Defining K6
sage: [(K.name, K.symmetry_type()) for K in K6]
[('K6_1', 'reversible'),
   ('K6_2', 'reversible'),
   ('K6_3', 'fully amphicheiral')]
```

three_genus()

Return the three genus of self.

From the KnotInfo description page:

The three-genus of a knot is defined to be the minimal genus of a Seifert surface for a knot.

To read the complete description type KnotInfo.K0_1.items.three_genus.description_webpage().

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.K5_2.three_genus() # optional - databsase_knotinfo
1
```

Note that this differs from the corresponding result in Sage since the latter is obtained for a Seifert surface that does not have the minimal genus:

```
sage: KnotInfo.K5_2.link().genus()
3
```

```
Bases: sage.structure.unique\_representation.UniqueRepresentation, sage.structure.sage\_object.SageObject
```

This class can be used to access knots and links via their index according to the series they belong to.

INPUT:

- crossing_number integer giving the crossing numer of this series of links
- is_knot boolean whether this series is a series of knots or proper links
- is_alternating boolean whether this series is restriced to alternating links or not This is not relevant for knots with less than 11 crossings
- name_unoriented string restricting the series to all links with that name_unoriented

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfoSeries
sage: K6 = KnotInfoSeries(6, True, True); K6
Series of knots K6
sage: K6(3)
<KnotInfo.K6_3: '6_3'>
sage: list(K6)
[<KnotInfo.K6_1: '6_1'>, <KnotInfo.K6_2: '6_2'>, <KnotInfo.K6_3: '6_3'>]
```

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```
sage: L6a = KnotInfoSeries(6, False, True); L6a
Series of links L6a
sage: L6a(2)
Series of links L6a2
sage: _.inject()
Defining L6a2
sage: list(L6a2)
[<KnotInfo.L6a2_0: 'L6a2{0}'>, <KnotInfo.L6a2_1: 'L6a2{1}'>]
sage: L6a2(0).series() == L6a
True
sage: L6a2(0) == L6a2('0')
True
```

inject(verbose=True)

Inject self with its name into the namespace of the Python code from which this function is called.

INPUT:

• verbose – boolean (optional, default True) to suppress the message printed on the invocation

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfoSeries
sage: KnotInfoSeries(6, True, True).inject()
Defining K6
sage: K6(2)
<KnotInfo.K6_2: '6_2'>
```

is_recoverable(unique=True, max_samples=8)

Return if all items of self can be recovered from its conversion to Sage links using the pd_notation and the braid_notation and their mirror images.

The method is indirectly used by the TestSuite.

INPUT:

- unique boolean (optional, default=``True``) see KnotInfoBase.is_recoverable()
- max_samples non negative integer or infinity (optional, default 8) limits the number of items to check (random sample). If set to infinity then no limit is set.

```
sage: from sage.knots.knotinfo import KnotInfo
sage: KnotInfo.L4a1_0.series().inject()
Defining L4a
sage: L4a.is_recoverable()
True
sage: L4a.is_recoverable(unique=False)
True
sage: KnotInfo.L5a1_0.series().inject()
Defining L5a
sage: L5a.is_recoverable()
False
sage: L5a.is_recoverable(unique=False)
True
```

list(*oriented=False*, *comp=None*, *det=None*, *homfly=None*)
Return this series as a Python list.

INPUT:

- oriented boolean (optional, default False) it only affects series of proper links. By default the list items of a series of proper links are again series of links collecting all orientation types of an unoriented name. To obtain the list of the individual links this keyword has to be set to True
- comp (optional, default None) if given an integer for this keyword the list is restriced to links having the according number of components. This keyword implies oriented=True
- det (optional, default None) if given an integer for this keyword the list is restriced to links having the according value for its determinant. This keyword implies oriented=True
- homfly (optional, default None) if given a HOMFLY-PT polynomial having normalization='vz' for this keyword the list is restriced to links having the according value for its HOMFLY-PT polynomial. This keyword implies oriented=True

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfoSeries
sage: K6 = KnotInfoSeries(6, True, True); K6
Series of knots K6
sage: K6.list()
[<KnotInfo.K6_1: '6_1'>, <KnotInfo.K6_2: '6_2'>, <KnotInfo.K6_3: '6_3'>]
sage: KnotInfoSeries(2, False, True).inject()
Defining L2a
sage: L2a.list()
[Series of links L2a1]
sage: L2a.list(oriented=True)
[<KnotInfo.L2a1_0: 'L2a1{0}'>, <KnotInfo.L2a1_1: 'L2a1{1}'>]
```

lower_list(oriented=False, comp=None, det=None, homfly=None)

Return this series together with all series with smaller crossing number as a Python list.

INPUT:

- oriented boolean (optional, default False) see the description for list()
- comp (optional, default None) see the description for list()
- det (optional, default None) see the description for list()
- homfly (optional, default None) see the description for list()

EXAMPLES:

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sage.knots.knotinfo.eval_knotinfo(string, locals={}, to_tuple=True)

Preparse a string from the KnotInfo database and evaluate it by sage_eval.

INPUT:

- string string that gives a value of some database entry
- locals dictionary of locals passed to sage_eval

EXAMPLES:

```
sage: from sage.knots.knotinfo import KnotInfo, eval_knotinfo
sage: L = KnotInfo.L4a1_0
sage: L.braid_notation(original=True)
'{4, {1, -2, 3, -2, -1, -2, -3, -2}}'
sage: eval_knotinfo(_)
(4, (1, -2, 3, -2, -1, -2, -3, -2))
```

sage.knots.knotinfo.knotinfo_bool(string)

Preparse a string from the KnotInfo database representing a boolean.

INPUT:

• string – string that gives a value of some database entry

```
sage: from sage.knots.knotinfo import knotinfo_bool
sage: knotinfo_bool('Y')
True
```

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