Restriction Sites Algorithm Element

The *element* searches restriction sites in the input sequence.

Parameters in GUI

Parameter	Description	Default value
Annotate As	Name of the result annotations.	Special value <rsite> is used. It specifies to use the enzymes names as names of the result annotations.
Circular	If *True* allows to search for restriction sites between the end and the beginning of the sequence.	False
Enzymes	Restriction enzymes used to recognize the restriction sites.	You must specify a value!

Parameters in Schema File

Type: rsite

Parameter	Parameter in the GUI	Type
key	**Annotate As**	*string*
circular	**Circular**	*boolean*
enzymes	**Enzymes**	*string*