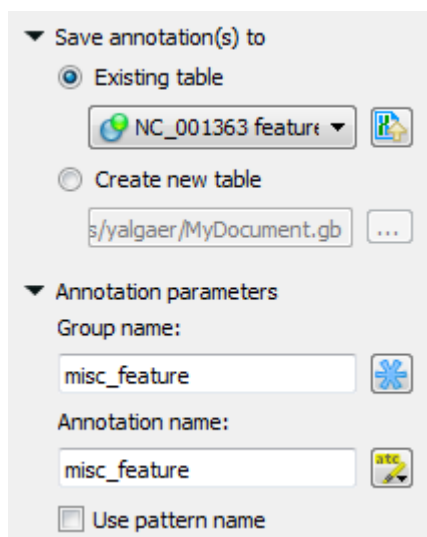


Annotations Settings



The screenshot shows a dialog box titled "Annotations Settings". It has two main sections. The first section, "Save annotation(s) to", has two radio buttons: "Existing table" (selected) and "Create new table". Under "Existing table", there is a dropdown menu showing "NC_001363 feature" and a file icon. Under "Create new table", there is a text field containing "s/yalgaer/MyDocument.gb" and a file icon. The second section, "Annotation parameters", has two text fields: "Group name:" with "misc_feature" and "Annotation name:" with "misc_feature". There are icons next to these fields. At the bottom, there is a checkbox labeled "Use pattern name" which is unchecked.

In the *Save annotation(s) to* group of parameters you can set up a file to store annotations. It could be either an existing annotation table object or a new document (file).

In the *Annotation parameters* group you can specify

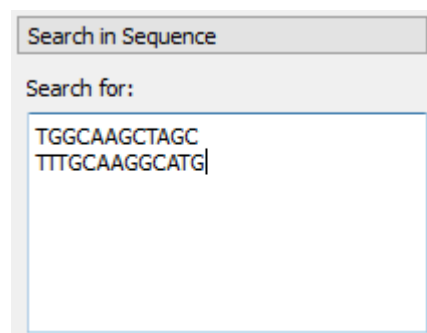
- A group that the found annotations will be stored in (this affects the name of the folder in the [Annotations Editor](#))
- The way name(s) of the found annotations are assigned (see below)

Searching for one or several patterns and names of the result annotations

If you search for one pattern only, than input the required name into the *Annotation name* field and leave the *Use pattern name* check box unchecked.

You can also search for several patterns at a time by:

- Inputting several patterns into the search field (click *<Ctrl>* + *<Enter>* keys to insert to a new line):



The screenshot shows a dialog box titled "Search in Sequence". It has a text field labeled "Search for:" containing two lines of text: "TGGCAAGCTAGC" and "TTTGAAGGCATG".

- Inputting several patterns into the search field in FASTA format:

Search in Sequence

Search for:

```

>pattern1
TGGCAAGCTAGC

>pattern2
TTTGCAAGGCATG

```

- [Loading patterns from a FASTA file](#)

Even when you search for several patterns, names of the found annotations will be identical by default (the name is specified in the *Annotation name* field).

If you want to assign different names to annotations found for different patterns, than you should:

- Input the patterns in FASTA format (the latter two cases above)
- Check the *Use pattern name* checkbox in the *Annotation parameters* group

Here is an example of the found annotations in the [Annotations Editor](#):

Annotations [MyDocument_1.gb] *	
misc_feature (0, 4)	
▶ pattern1	24..35
▶ pattern1	5267..5278
▶ pattern2	50..62
▶ pattern2	5293..5305