

Captions on Annotations

It is possible to show a value of a qualifier of an annotation instead of the annotation type name in the *Sequence Zoom View*. To enable this option for an annotation type check the *Show value of qualifier* check box and input the values of the required qualifiers in the text field nearby this check box. See the image below.

The screenshot displays a sequence viewer for NC_001363. The top part shows a genomic map with a blue arrow indicating a 589 bp region. Below this, the DNA sequence is shown with a yellow highlight over a specific region. The bottom part of the interface shows a tree view of features and a configuration panel on the right.

Name	Type	Value
NC_001363 features [m...]		
source (0, 1)		
misc_feature (0, 2)		
misc_feature	Misc. Feature	5245..5833
note		Show value of qualifier
misc_feature	Misc. Feature	2..590
comment (0, 1)		
CDS (0, 4)		
CDS	CDS	5048..5203
CDS	CDS	3875..4999

Annotations Highlighting

Select an annotation name:

Annotation	Color
BadI	Light Green
BaeGI	Orange
BaeI	Purple
CDS	Green
SacI	Cyan
comment	Pink
misc_feature	Yellow
source	Grey

Show all annotation names

Configure the annotations:

- Show annotations
- Show on translation
- Show value of qualifier:

note

Help

If you input several qualifiers names (separated by comma), then the first found qualifier is taken into account and shown on the annotation.