

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 the NC_001363 [d] sequence viewer. The top panel shows a genomic map with a scale from 1 to 5k. A red arrow points to a blue bar representing a feature, with the text "Show value of qualifier" above it. Below the map, the DNA sequence is shown with various annotations. A red arrow points to a specific annotation in the list below the sequence.

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

The right panel shows the "Annotations Highlighting" settings. Under "Configure the annotations:", the "Show value of qualifier" checkbox is checked. A red arrow points to this checkbox. Below the checkbox is a text field labeled "note".

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