

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 bioinformatics software interface with a sequence viewer and a configuration panel on the right.

Sequence Viewer:

- Top track: NC_001363 [d]
- Scale: 1 to 5k (500, 1k, 1.5k, 2k, 2.5k, 3k, 3.5k, 4k, 4.5k, 5k)
- Annotation: **Show value of qualifier** (orange bar)
- Sequence: 5 245 5.3k 5350 5.4k 5450 5.5k 5550 5.6k 5650 5.7k 5750 5 833
- Sequence: A K Q D I C G K Q F L P R L
- Sequence: G P N R I S V V S S C P G S
- Sequence: G Q T G Y L W * A V P A P A R
- Sequence: GGCCTAACAGGATATCTGTGGTAAGCAGTTCCTGCCCGGGCTCG
- Sequence: 5 445 5450 5455 5460 5465 5470 5475 5480 5 488
- Sequence: CCGGTTTGTCTATAGACACCATTCGTCAAGGACGGGGCCGAGC
- Sequence: P W V P Y R H Y A T G A G A R
- Sequence: G F L I D T T L L E Q G P E
- Sequence: A L C S I Q P L C N R G R S P

Annotations Panel:

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 Panel:

Select an annotation name:

Annotation	Color
BadI	Light Green
BaeGI	Orange
BaeI	Purple
CDS	Green
SacII	Light Blue
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.