

Highlighting Annotations

To configure settings of annotation names go to the *Annotation Highlighting* tab in the *Options Panel*.

By default the tab shows annotations names of the opened *Sequence View*.

The screenshot displays a sequence viewer for NC_00136. The top part shows a scale from 1 to 5,833 bp. Below the scale, a sequence is shown with various annotations. A specific region from 3875 to 4999 bp is highlighted in blue, with a length of 1125 bp. The sequence is shown in three lines: the first line has amino acid codes (D H P L D * H G A F N A M), the second line has nucleotide codes (T I L * T D M A H S T P C), and the third line has the full nucleotide sequence (GACCATCCTCTAGACTGACATGGCGCATTCAACGCCATGCTGGTAGGAGATCTGACTGTACCGCGTAAGTTGCGGTACGDELSVH R M * R W A V M R * V S M A C E V G H S W G R S Q C P A N L A M). Below the sequence, a list of annotations is shown with columns for Name and Type. The annotations include CDS, comment, misc_feature, and source. To the right, the 'Annotations Highlighting' panel is open, showing a table of annotation names and their colors. The table lists: BadI (light green), BaeGI (orange), BaeI (purple), CDS (green), SacII (cyan), comment (pink), misc_feature (tan), and source (grey). Below the table, there are checkboxes for 'Show annotations', 'Show on translation', and 'Show value of qualifier'. The 'Show value of qualifier' checkbox is checked, and a text box contains 'label,note'. There are also navigation buttons (Previous and Next) and a 'Help' button.

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

If you want to see all annotation names, click the *Show all annotation names* link. The *Previous annotation* and *Next annotation* buttons seek to the previous or to the next annotation of the view correspondingly.

Find below information about annotations names' properties that you can configure.

- [Annotations Color](#)
- [Annotations Visibility](#)
- [Show on Translation](#)
- [Captions on Annotations](#)