

Highlighting Annotations

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

By default the tab shows annotations types of the opened Sequence View.

The screenshot shows the BioEdit software interface with the 'Annotations Highlighting' tab selected in the Options Panel. The main window displays a sequence view for NC_001363 [dna]. The sequence is shown with various annotations highlighted in different colors: orange for CDS, green for misc_feature, and dark green for source. A detailed view of the sequence highlights specific codons and features. Below the sequence, a table lists the names and values of the annotations:

| Name | Value |
|--|-----------------------------|
| Auto-annotations [murine.gb NC_001363] | |
| NC_001363 features [murine.gb] | |
| CDS (0, 4) | |
| CDS | 1042..2658 |
| CDS | join(2970..3413,3412..3873) |
| CDS | 3875..4999 |
| CDS | 5048..5203 |
| misc_feature (0, 2) | |
| misc_feature | 2.590 |
| misc_feature | 5245..5833 |

The right side of the interface includes a legend for annotation types (CDS, misc_feature, source), configuration options for each type (Show annotations of this type, Show on translation, Show value of qualifier), and navigation buttons for previous and next annotations.

If you want to see all annotation types, click the *Show all annotation types* 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 types' properties that you can configure.

- Annotations Color
- Annotations Visibility
- Show on Translation
- Captions on Annotations