

# Details View

The *Sequence details view* is a supplementary component of the *Sequence overview*. It is used to show sequence content without zooming. Every time you double click the sequence in the *Sequence overview* area or select an annotation, the corresponding sequence position is made visible in the *Sequence details view*.

For a DNA sequence the *Sequence details view* automatically shows complement DNA strand and 6 amino translation frames.

The screenshot shows the UGENE software interface with the following components:

- Project Panel:** Shows the sequence NC\_001363 [dna] with a scale from 1 to 5.833 kb.
- Annotation Panel:** Shows a CDS annotation from 2970.31k to 5017.1k, with a selected region from 3875 to 4999 (1125 bp).
- Sequence View:** Displays the original sequence (top), complement strand (middle), and six amino acid translations (bottom).
- Annotations:** Includes 'Sacl (1)', 'misc\_feature (2)', and a 'Genetic code selector'.
- Table:** A table listing CDS annotations with their coordinates.

Name	Type	Value
CDS	CDS	join(2970..3413,3412..3873)
CDS	CDS	3875..4999
CDS	CDS	5048..5203
comment (0, 1)		
misc_feature (0, 2)		

See also:

- [Navigating the Details View using the Sequence Overview](#)
- [Translating Nucleotide Sequence](#)