

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.
- Annotations:** NP 597742.2 and NP 040336.1 are visible as orange bars. A specific CDS annotation is selected, spanning from 3875 to 4999 bp.
- Sequence View:** Displays the original DNA sequence (top) and its complement (bottom). The selected CDS region is highlighted in yellow.
- Amino Acid Translations:** Six amino acid sequences are shown. The top three are labeled "3 direct amino translations" and the bottom three are labeled "3 complement amino translations".
- Genetic Code Selector:** A dropdown menu is set to "C" (Standard Genetic Code).
- Table:** A table at the bottom lists annotations. The selected CDS entry is highlighted.

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](#)