

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 Sequence details view for a DNA sequence (NC_001363). The view displays the original sequence, its complement, and three amino acid translations for each. Annotations are visible at the top, and a table at the bottom lists the selected annotation.

Annotations:

- NP 597742.2
- NP 597742.2
- NP 040336.1
- SacI (1)
- misc_feature (2)

Sequence details:

29703.1k 3.2k 3.3k 3.4k 3.5k 3.6k 3.7k 3.8k 3.9k 4k 4.1k 4.2k 4.3k 4.4k 4.5k 4.6k 4.7k 4.8k 5 017

3875 [1125 bp]

Annotations:

- Toggle visibility
- 3 direct amino translations
- The original sequence
- Genetic code selector
- The complement strand
- 3 complement amino translations
- The selected annotation

Table:

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](#)