

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:

Original sequence: E Q L D H P L D *

Complement strand: M A H S T P C S Q T

3 direct amino translations: E Q L D H P L D *

3 complement amino translations: M A H S T P C S Q T

Genetic code selector: The genetic code selector is set to the standard genetic code.

The selected annotation: The selected annotation is the CDS (Coding Sequence) annotation.

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](#)