

Sequence 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 the NC_001363 sequence. The top toolbar includes a 'Genetic code selector' and a 'Toggle visibility' button. The main window displays a DNA sequence with annotations for CDS (NP_040335.1, NP_59774, NP_040336.1) and genes (Msvgp1, Msvgp2, Msvgp3). Red arrows point to various features: 'Genetic code selector' and 'Toggle visibility' in the top toolbar; '3 direct amino translations' and 'The original sequence' pointing to the top DNA strand; 'The complement strand' pointing to the bottom DNA strand; 'The selected annotation' pointing to the 'CDS' feature in the left sidebar; and '3 complement amino translations' pointing to the bottom DNA strand. The bottom sidebar shows a list of features with their coordinates.

Name	Value
NC_001363 features [NC_001363.gb]	
CDS (0, 4)	
CDS	1042..2658
CDS	join(2970..3413,3412..3873)
CDS	3875..4999
CDS	5048..5203
gene (0, 4)	

See also:

- [Navigating the Sequence details view using the Sequence overview](#)
- [Selecting Amino Translation](#)
- [Showing and Hiding Translations](#)