# **UGENE** Terminology

# Project

Storage for a set of data files and visualization options.

# Document

A single file (can be stored on a local hard drive or be a remote web page). Each document contains a set of objects.

# Object

A minimal and complete model of biological data. For example: a single sequence, a set of annotations, a multiple sequence alignment.

#### Task

A process, usually asynchronous, that works in background. For example: some computations, loading and writing files.

## Plugin

A dynamically loaded module that adds new functionality to UGENE.

#### **Object View**

A graphical view for a single or a set of objects.

#### **Project View**

A visual component used to manage active project.

#### **Task View**

A visual component used to manage active tasks.

# Log View

A visual component used to show logs.

#### Notifications

A visual component used to show notifications. Generally it is used to open tasks reports.

# **Plugin Viewer**

A visual component used to manage plugins.

#### Sequence View

An Object View aimed to visualize DNA, RNA or protein sequences along with their properties like annotations, chromatograms, 3D models, statistical data, etc.

#### Annotation

Additional information about a sequence, identified by its name and the sequence region.

#### **Alignment Editor**

An Object View used to visualize and edit DNA, RNA or protein multiple sequence alignments.

#### **Options Panel**

An Options Panel it is the panel with different information tabs and tabs with settings for Sequence View and Assembly Browser.

In the image below you can see a typical UGENE window with a Project View and a single Object View window opened:

