Importing APR and ACE Files

To start working with APR file you can open it in the Alignment Editor in read-only mode or convert it to another format.

To do this, open the *.apr file. The following dialog will appear:

Select Document Format	3		
Options for gyrA.apr			
Open in read-only mode			
 Convert to another format: 			
Save to fiie ugene/data/samples/APR/gyrA			
File format PHYLIP Sequential			
Help Scancel VC]		

If you choose the first option the file will be opened in the Alignment Editor as multiple sequence alignment in read-only mode. If you choose the second option the file will be converted to a corresponding format and open in UGENE.

The following formats are available: CLUSTALW, FASTA, Mega, MSF, NEXUS, Phylip Interleaved, Phylip Sequential, Stockholm.

To start working with ACE file you can open it in the Alignment Editor or import it to the UGENE database file.

To do this, open the *.ace file. The following dialog will appear:

Select Docum	nent Format	8	
Options for K26.	ace		
Multiple set	quence alignment in the Alignment Edito	r	
 Short reads assembly in the Assembly Browser 			
Save to file	lata/samples/ACE/K26.ace.ugenedb		
Help	<mark>⊗ C</mark> ancel √ <u>O</u> K		

If you choose the first option the file will be opened in the Alignment Editor as multiple sequence alignment. If you choose the second option the file will be opened in Assembly Browser.