Sanger Reads Editor Features

Sanger Reads Editor is dedicated to DNA sequence analysis and manipulation. You can trim, map to reference and view Sanger sequencing trace files and create consensus sequences.

The editor provides interactive visual representation which includes:

- Navigation through an chromatogram;
- Flexible zooming;
- Several consensus calculation algorithms.

Using the *Sanger Reads Editor* you can:

- Export alignment without chromatogram;
- Edit chromatogram: remove/replace/trim.