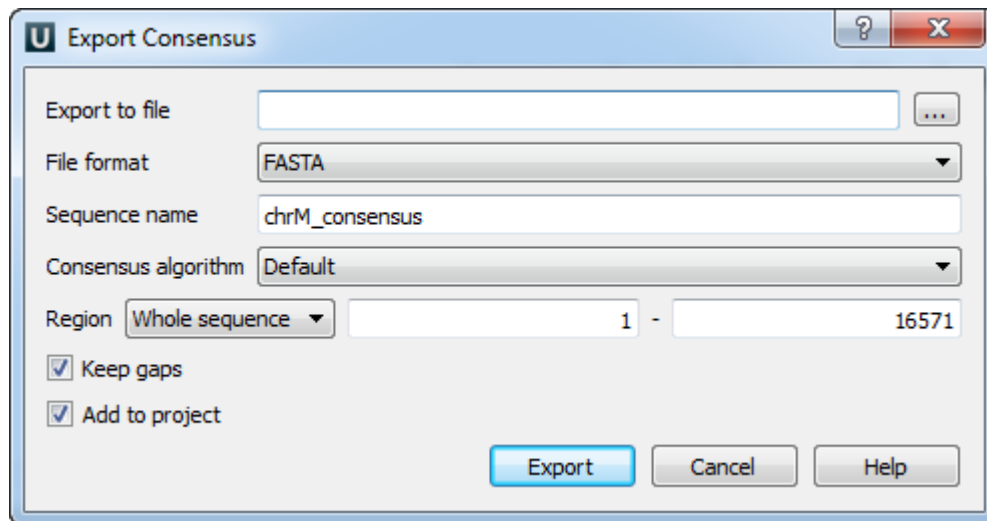


Exporting Consensus

To export a consensus sequence of the assembly, select either the *Export consensus* item in the *Consensus Area* context menu or the *Export Consensus* item in the *Reads Area* context menu.

The *Export Consensus* dialog appears:



The screenshot shows the 'Export Consensus' dialog box. It has a title bar with a question mark icon and a close button (X). The dialog contains the following fields and controls:

- Export to file:** A text input field with a browse button (three dots) to its right.
- File format:** A dropdown menu currently showing 'FASTA'.
- Sequence name:** A text input field containing 'chrM_consensus'.
- Consensus algorithm:** A dropdown menu currently showing 'Default'.
- Region:** A dropdown menu currently showing 'Whole sequence', followed by two input fields for coordinates: '1' and '16571'.
- Keep gaps:** A checked checkbox.
- Add to project:** A checked checkbox.
- Buttons:** 'Export', 'Cancel', and 'Help' buttons at the bottom right.

Select a file and the file format. The consensus can be exported to a FASTA, FASTQ, GFF or GenBank file.

Modify, if required, the exported sequence name and choose the [consensus algorithm](#).

The consensus is exported with gaps if the *Keep gaps* check box has been checked.

Also you can select the exporting region. It can be either a *Whole sequence*, a *Visible* region, or a *Custom* region.

When all the parameters are set click the *Export* button.

The consensus sequence is exported to the file and if the *Add to project* check box has been checked it is added to the current *project* and opened.