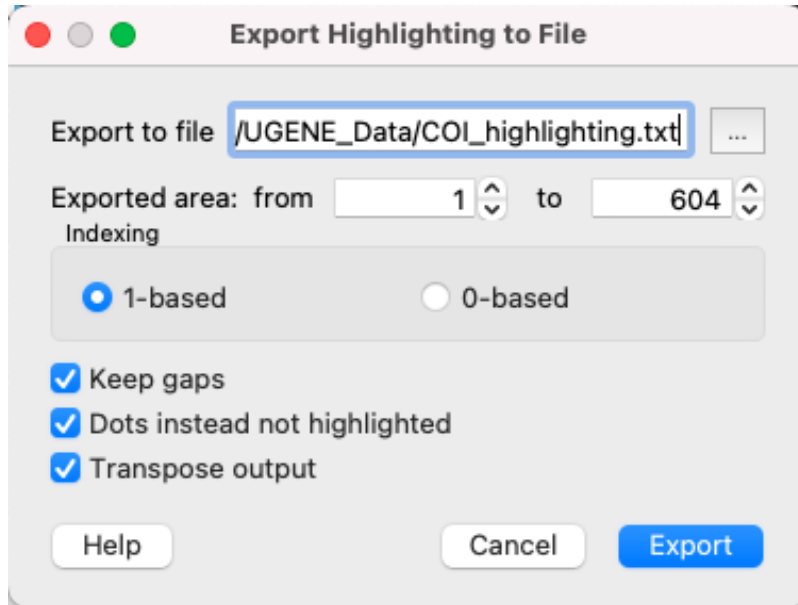# Export Highlighting to File

Select the output file to export the highlighting, choose exported area and click on the *Export* button. The task report will appear in the *Notifications*.

Also, you may set additional parameters:

- **Indexing** - select the number to start counting from:
    - *1-based* - the first column has number **1**,
    - *0-based* - the first column has number **0**.
- **Keep gaps** - check if you want to store gap symbols in the result file.
- **Dots instead not highlighted** - replace not-highlighted bases with dots in the result file.
- **Transpose output** - if this checkbox is checked, the sequences will be located horizontally (not vertically, as in UGENE). May be useful, if you have a long alignment, but not many sequences.