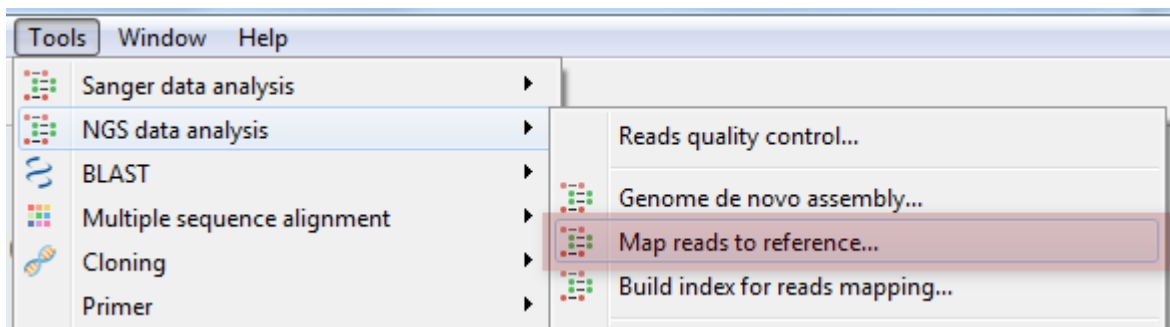


# Bowtie

*Bowtie* is a popular short read aligner. Click [this link](#) to open *Bowtie* homepage. *Bowtie* is embedded as an *external tool* into UGENE.

Open *Tools DNA Assembly* submenu of the main menu.



Select the *Align short reads* item to align short reads to a DNA sequence using *Bowtie*. Or select the *Build index* item to build an index for a DNA sequence which can be used to optimize aligning of the short reads to the sequence.

- [Bowtie Aligning Short Reads](#)
- [Building Index for Bowtie](#)