



Ready-to-Sequence sample requirements

"Ready to sequence" samples are a library or pool of libraries sent for sequencing only. The TGC unit will not be held responsible for sequencing bias or low complexity regions due to self-made libraries.

- Please submit a <u>pooled</u> library in a 1.7 ml tube labeled with PI name and date of submission.
- The final library pool should be at a minimum of 20 ul of a 4 nM pool.
- Please provide a Bioanalyzer or TapeStation results report (pdf) of your final library pool. If such a report is not provided, the TGC will perform the analysis for you at extra charge.
- An electronic Ready to Sequence Submission Form should be filled out and sent by email to linde@technion.ac.il. It is important to specify the following parameters:
 - o The samples' names and both indexes' sequence (not only name/number!).
 - The preferred loading concentration and % PhiX required.
 - Required read length.