# The Center for Genomic Technologies.

## Sample submission for Library construction for deep sequencing

**Please send the form by E-mail to: nadi.turjema@mail.huji.ac.il**

Date: _____

Sender (full name): _____     Head of the group: _____

Email address- _____     Phone_____

User ID at Basespace:_____

Samples characteristic: _____     Desired Library type: _____

Sequencing machine:_____     Sequencing length:_____

Number of sample to run together (for more than 12 samples):_____

Was sequencing kit ordered from Danyel Biotesch? _____

Remarks:_____

*For RNA, please submit 100ng-2ug (ideally 1ug) total RNA.
*For more than 4 samples please **submit in a strip/ strips or plate** and not in individual eppendorf. On the strip/plate write only the sample number corresponding to your sample name according to the table below. Samples in individual Eppendorf should have also sample number written on them according to the table below.
*We keep excess sample approximately a month after library construction, then we throw the samples without further notice.

Samples:

| Sample number | Sample name | Concentration | Volume ul | Total RNA ug |
|---|---|---|---|---|
| 1 | | | | |
| 2 | | | | |
| 3 | | | | |
| 4 | | | | |
| 5 | | | | |
| 6 | | | | |
| 7 | | | | |
| 8 | | | | |
| 9 | | | | |
| 10 | | | | |
| 11 | | | | |
| 12 | | | | |

Tel: 02-6585211     Fax: 02-6584048