**The Genomic Applications Lab**

**Sequencing project submission**

Date: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ PI: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

E-mail: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Phone: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

X

**Sample type:** DNA x RNA Library

Organism: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

**Library preparation protocol:**

XX

* V

X mRNAseq Total RNAseq (Ribosomal depletion) SMARTer pico

NexteraXT 16S Amplicon sequencing (2nd PCR)

Other: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

 *Kit used: \_\_\_\_\_\_Kapa Biosystems\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Company:\_\_\_\_\_\_\_\_\_\_\_\_*

**Run parameters:**

**Storage:**

No. of samples**:** \_\_\_\_\_

Box color: \_\_\_\_\_\_\_\_\_

New freezer drawer no.: \_\_\_\_

Platform: **x** NextSeq MiSeq

Reading length: Single Read: \_\_\_\_ bp

Paired end: \_\_\_\_\_ x \_\_\_\_\_ bp

V

Index: Single index: \_\_6\_\_\_ bp

 Dual index: \_\_\_\_\_ bp

**Samples:**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | Sample Name | Qbit ng/µl | Av. size | Index 1 | Index 2 | No. of reads  |
| 1 |  |  |  |  |  |  |
| 2 |  |  |  |  |  |  |
| 3 |  |  |  |  |  |  |
| 4 |  |  |  |  |  |  |
| 5 |  |  |  |  |  |  |
| 6 |  |  |  |  |  |  |
| 7 |  |  |  |  |  |  |
| 8 |  |  |  |  |  |  |
| 9 |  |  |  |  |  |  |
| 10 |  |  |  |  |  |  |
| 11 |  |  |  |  |  |  |
| 12 |  |  |  |  |  |  |
| Total amount of reads: |  |

 Index type: x TruSeq

 Lexogen

 Nextera Custom

*Sequencing date: \_\_\_\_\_\_\_\_\_\_*

\*\*\* **Long time storage** \*\*\*

Freezer: \_\_\_\_\_\_\_\_\_\_\_\_

Drawer: \_\_\_\_\_\_\_\_

Box color: \_\_\_\_\_\_\_\_\_\_\_

Comments:\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

V

SampleSheet sent by mail to: genomics@ekmd.huji.ac.il