 Illumina Genome Analyzer System Request Form

 **MIT BioMicro** <http://biomicro.mit.edu/>

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User Name:\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Email address\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Phone\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Date\_\_\_\_\_\_\_\_\_

Lab Name\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Account # / PO#\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Address\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

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| --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **Sample ID** | **Sample Type****(gDNA, smRNA…)** |  **Primer****(SR, PE, Custom)** | **uL** **(Total)** | **# bases (Forward)****(max = 36)** | **# bases (Reverse)****(max = 36)** | **Genome****(eg. Mm8)** | **BMC\_ID** |
| 1 |  |  |  |  |  |  |  |  |
| 2 |  |  |  |  |  |  |  |  |
| 3 |  |  |  |  |  |  |  |  |
| 4 |  |  |  |  |  |  |  |  |
| 5 |  |  |  |  |  |  |  |  |
| 6 |  |  |  |  |  |  |  |  |
| 7 |  |  |  |  |  |  |  |  |
| 8 |  |  |  |  |  |  |  |  |
| 9 |  |  |  |  |  |  |  |  |
| 10 |  |  |  |  |  |  |  |  |

**INCLUDE CONTROL LANE (circle one)**: YES | NO

\* Primer types include Single end DNA, Paired end DNA, and small RNA.

\*\* Sample concentration can be confirmed on the Agilent BioAnalyzer. Accurate concentration is important for cluster generation.

**BioMicro Use Only**

 **Unit Charge $\_\_\_\_\_\_\_ Total Charge back $\_\_\_\_\_\_\_\_\_\_ Database Tracking Number\_\_\_\_\_\_\_\_\_\_\_\_\_\_ DATE FINAL \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**