

Solexa // 454 Sequencing Request

User name	PI name
Contact e-mail	Center
Quotation num.	Date

Signature

	Sample name	Species	Sample type ¹	Read length	Conc. (ng/μl) ²	Vol. ₂ (μl)	ID
1							
2							
3							
4							
5							
6							
7							
8							
9							
10							
11							
12							
13							
14							

Bioinformatics Core will analyse your data: **YES** **NO**

- 1 ChIP Seq., Single-end Genomic DNA, Paired-end Genomic DNA, Mate-Pair, miRNA, smallRNA, mRNA Seq. or other applications
- 2 Indicate whenever possible. Please make sure to fit sample concentration and volume requirements. For detailed information click [here](#) for Solexa and contact the service in the case of 454 sequencing

Shaded columns are to be filled by the service

Users submitting samples with pre-assigned flow cell runs are requested to submit samples for library prep and quality check at least 3 weeks before their assigned run dates, otherwise runs will be rescheduled.

Information for users submitting libraries newly prepared by them:

- 1) Please submit libraries to the Unit at least 10 days ahead of the scheduled flow cell run. We will run the libraries on the Bioanalyzer, and will quantify the libraries (if necessary, with two different assays).
- 2) Failure to provide the libraries 10 days in advance will result in your run being rescheduled.
- 3) Only libraries you definitely want to run should be submitted to the service.