

## Experiment information

### Content

Please request the following information from the sequencing lab for each sequencing experiment:

**experiment\_id:** This is an internal id for your experiment record. It must be unique to one experiment record for a provider. Example is 'e123'. The value must be a single word.

**instrument\_model:** Example is Illumina.

**instrument\_model:** Example is Illumina MiSeq

**library\_name:** Any library name.

**library\_layout:** PAIRED/ SINGLE

**library\_strategy:** Example is WGS

**library\_source:** Example is GENOMIC

**library\_selection:** Example is RANDOM

**insert\_size:** For paired end reads only. Example is 250.

### Format

The accepted format is tab-delimited file. The first line shall contain the list of keys. From the second line the values for corresponding keys need to be provided. Each line shall represent one experiment record.

A complete example has been provided in **experiment.txt** file.