



DNA / RNA SEQUENCING CONTACT (NIG)

Date:

Project number (filled by NIG) :

Project title (with one sentence) :

Project Description :

CONTACT INFORMATION

Contact person

Name :

Email :

Address :

Phone :

Group leader/Principal investigator

Name :

Email :

Address :

Phone :

PROJECT INFORMATION AND/OR PROTOCOL INFORMATION

Library type

single-end library

paired-end library

Read length

☐ Illumina

☐ Illumina

☐ MiSeq

☐ Minion

☐ 30 bp

☐ 100 bp

☐ 150 bp

☐ 200 bp

Analysis application type

total RNA-seq

small RNA-seq

ChIP-seq

Protein sequencing

de novo genome sequencing

total RNA-seq

small RNA-seq

ChIP-seq

Targeted DNA-seq

genome resequencing

ATAC-seq

Amplification-seq

No RNA

Sample pooling

Yes

No

if yes, how many samples will be pooled together? _____

Total number of samples (including biological replicates) :

Organism:

Sample type :

DNA / RNA extraction method :

Other possible pre-processing done to the samples (please specify all applied protocols)

Disease treatment: