



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 / SEQ. PROTOCOL INFORMATION

Library type

single-end library

paired-end library

Read length

☐ 50bp

☐ Illumina200

☐ 100bp

☐ Illumina400

☐ 150bp

☐ MiSeq

☐ 200bp

☐ Minion

Analysis application type

total RNA-seq

smallRNA-seq

ChIP-seq

Resolute sequencing

de novo genome sequencing

total RNA-seq

smallRNA-seq

RRBS

Targeted DNA-seq

genome resequencing

ATAC-seq

Amplicon-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: