

## HOW TO REQUEST A SEQUENCING SERVICE TO **CRG GENOMICS UNIT**:

1. Please, **send an e-mail** to [ngs@crg.eu](mailto:ngs@crg.eu) with the following information:
  - Number of samples
  - Type of service (application)
  - How to sequence (Single Read or Paired End, length of the read)
  - Number of lanes needed or number of million reads per sample
  - Information needed for the quotation of the service
    - o Name of the PI (always)
    - o Name of the Institute (always)
    - o Address (for new users)
    - o VAT number of the institute (for new users)

If you do not know how to sequence and require information and advice, please, explain briefly your project and what do you want to analyze so we can discuss with you the best strategy for you.

2. **Wait for the reply from the Unit** with the instructions and the approval to send the samples. Please, do not send / bring any sample without the approval from Genomics Unit.
3. Once you have the approval from the Unit, please **send an e-mail** to [ngs@crg.eu](mailto:ngs@crg.eu) with the filled "**Sequencing Request Form**"

**IMPORTANT:** Subject of the mail should include:

- **QC service:** if it is for a sample quality control service
- **Sequencing project:** if it is for a sequencing service
- **Microarrays project:** if it is for a microarrays service



**cnag**

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