

POLICY GUIDELINES

The CNAG operates according the following **policy guidelines**:

1. The CNAG provides assistance to researchers, primarily in the field of genomics, by offering expertise in project design, sequencing, data analysis and interpretation.
2. The CNAG sequencing services are open to researchers from the Spanish, European and International Research Community, both for academic research and industrial applications.
3. The CNAG sequencing services are provided on a fee-for-service basis. Non-academic users are charged the full-cost fee. However, sequencing costs for academic institutions and other public sector research organizations are subsidized to reflect the public investment made to enhance the accessibility of the facility.

For more information on the CNAG sequencing platform and its applications, visit www.cnag.eu.

COMMITMENTS

Users of CNAG are expected to adhere to the following **commitments**:

1. Users must take responsibility for declaring that the samples were obtained with the necessary approval from the Bioethics Committee of the Depositor. In the case of human samples, users must declare that they have been obtained with informed consent from the donors, both for collection and their use in the proposed experiments.
2. Users are required to acknowledge the work carried out by the CNAG in any or all of the following ways:
 - a) If the CNAG has made intellectual contributions, collaborators should engage in discussions with the relevant CNAG scientists or staff to determine appropriate co-authorship for potential and pending publications based on these contributions.
 - b) At the very least, peer-reviewed publications must acknowledge the work of the CNAG. This acknowledgment should be included either in the Acknowledgment or Materials and Methods sections.
3. Users are expected to authorize CNAG to cite its identifiable information in its outreach or scientific activities and to provide additional information and documentation for use as dissemination material at CNAG.

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Written by:	Review by:	Approved by:	Date:

ACCESS

Two access modalities are considered, depending on the amount of sequencing required:

1. Access for small-medium scale projects

- a) Researchers should contact the CNAG project manager team (projectmanager@cnag.eu) to discuss the project feasibility, design and associated costs.
- b) The activity call is continuously open, allowing researchers to submit their proposals at any time.
- c) All experiments receive the same priority.

2. Access for large scale projects

- a) Researchers must submit an application for reviewed. Application forms for proposals can be found at [Access | Omicstech](#)
- b) Applications are reviewed on a semestral basis by an Access Committee consisting of four members, including the CNAG director and three Genomics Experts.
- c) Accepted proposals are published, while explanations are provided in confidence for those not accepted.
- d) The evaluation of proposals is based on the following principles and criteria:
 - ✓ The quality and relevance of the overall scientific project, including the proposed activities (50%).
 - ✓ Evidence of strong institutional support and funding for the lead applicant (25%).
 - ✓ Scientific credentials of the applicant's research group (25%).
- e) Consultation with CNAG personnel BEFORE submitting a grant application involving large-scale sequencing is necessary.
- f) Large-scale projects are defined as follows:
 - ✓ Whole genome sequencing (WGS) human projects that include more than 50 samples
 - ✓ Whole exome sequencing (WES) and RNA-seq projects that include more than 100 samples
 - ✓ *de novo* assembly and annotation projects that include more than 5 samples
 - ✓ Single Cell analysis projects that include more than 30 samples
 - ✓ Multi-application projects: projects that require different applications and technologies

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