

Public call for collaborative research and research meeting for FY 2026
The Medical Institute of Bioregulation, Kyushu University,
(Start of the public call: December 2025)

Having been recognized as a Joint Usage/Research Center for the Multi-Stratified Host Defense System since April 1, 2010, the Medical Institute of Bioregulation (MIB) , Kyushu University has been promoting collaborative research with researchers involved in host-defense studies.

We publicly call for research proposals for fiscal year 2026

1. Research Categories for which a Call for Proposals is Organized

A. Collaborative research

(1) Instrument usage type project:

Research project using the multi-stratified levels of research facility infrastructure and the technology of this institute. Research expenses are distributed to cover the instrument usage fee, travel expenses and so on.

(2) Collaborative research type project:

Research project aiming at clarifying the host-defense system using the knowledge about host defense accumulated in this institute. Only travel and accommodation expenses are granted.

B. Research meetings

Meetings focusing on information exchange and presentation of research results among researchers involved in host-defense studies, or meetings for planning collaborative research among such researchers.

2. Research Field

We call for collaborative research proposals on the following four research fields. The available facilities and research support service for each field are listed (Attachment). For information on facilitators, please see the following link (Laboratory List):

https://www.bioreg.kyushu-u.ac.jp/mib/labo_list_e.html

(1) Genome Science (including Genomics, Epigenomics, Transcriptomics, and Informatics Analysis)
Genomics:

We provide services of massive detection of genomic variations. We accept collaborative researches such as large-scale genome sequencing using NovaSeq or HiSeq. We also accept mid-to-small scale researches such as exome sequencing of disease pedigrees and Amlicon-seq of biomes using MiSeq. Furthermore, starting from fiscal year 2024, we will support the determination of structural polymorphism in model organisms such as humans and mice, sequence acquisition for non-model organisms, and metagenomic sequencing using the PacBio Sequel IIe long-read sequencer, which provides high accuracy and longer reads compared to the short-read NovaSeq.

Epigenomics:

We plan to perform collaborative research on epigenetic modifications, such as DNA methylation and histone modifications, which are important for the maintenance of function and homeostasis of living tissues. For example, using our cutting-edge technology applicable to a limited amount of DNA, we are able to reveal the genome-wide distribution pattern of DNA methylation at single-base resolution. Such studies will provide a comprehensive understanding of epigenetics and contribute to the progress of biosciences related to health and diseases.

Transcriptomics:

We accept proposals for collaborative studies on understanding cell homeostasis and its deficiency appearing as abnormal development, tumorigenesis, and ageing using transcriptome quantification and/or epigenome regulation. Single Cell transcriptomics has enabled elucidation of the phenotype of an individual not only by identifying the cells responsible for the phenotype but also by understanding the cell dynamics reconstructed from capturing cell population. Applicants are encouraged to apply these comprehensive analyses for humans or other model/non-model organisms with appropriate statistical analyses.

Information Analysis Infrastructure:

We have established an analysis system for sequence data, with a focus on genomics, which can be used primarily to analyze data generated through collaborative research. In addition, we have developed an international database of human haplotypes—the Joint Open Genome and Omics Platform (JoGo) (<https://jogo.csml.org>)—and we welcome proposals for collaborative research that leverage both this haplotype database infrastructure and our analysis infrastructure.

(2) Spatial Omics (including Spatial transcriptomics, Spatial proteomics)

Spatial omics technologies enable the quantification of the localization of numerous molecules while preserving the spatial organization of molecules and cells within tissues or intracellular environments. Collaborative research is conducted utilizing spatial multi-omics approaches, including transcriptome analysis in specific tissue regions or intracellular areas using the Photo-Isolation Chemistry method, proteome analysis through sequential immunostaining, and spatial multi-omics in cellular and tissue samples using sequential RNA/DNA-FISH techniques.

(3) Structural Biology

We plan to conduct collaborative research for the determination of protein structures using structural biology technologies, such as cryo-electron microscopy (cryo-EM) and X-ray crystallography.

The high-resolution structure analysis provides atomic- and molecular-level insights into how proteins work to fulfill their functions, and allows drug design that specifically controls protein functions. Notably, cryo-EM analysis sometimes enables determination of protein structures that are hard to be solved by X-ray crystallography, such as membrane proteins and biomacromolecule complexes.

(4) Embryonic and Genetic Engineering

We plan to conduct collaborative research on the elucidation of host-defense mechanism at an individual level by developmental engineering methods using mice, and the development of new therapies for diseases caused by the breakdown of the host-defense mechanism. Generation of genetically engineered mice by developmental engineering includes the production of knockout, knock-in, and transgenic mice from fertilized eggs and ES cells. We also actively introduce genome editing technology based on the CRISPR/Cas9 system.

3. Eligibility for Applicants

Researchers belong to universities, national or public institutions, or equivalent institutions

4. Period of Research

From April 1, 2026 to March 31, 2027

5. How to Apply

Please fill out the application form and send it to the address (see 11. Contact below) by post or e-mail. You MUST discuss the details of the collaboration with the facilitator of the MIB, before sending your completed application form.

The application form is downloadable at: https://www.bioreg.kyushu-u.ac.jp/mib/activities_collabo_e.html

Required Documents:

- A. Collaborative research : application form 1
- B. Research meeting : application form 2

Application deadline:

February 25, 2026

* Each research group is not allowed to apply two or more collaborative research projects per year, and the same application title must not be used more than 2 years.

*We may accept your application even after the application deadline. Please contact the facilitator of the MIB or the contact person (see “11. Contact” below) in advance. In case that we stop accepting applications, we will notify in the MIB HP.

6. Screening Results

Once the review committee has made a decision to either accept or reject the application, applicants will be informed of the result as soon as possible.

7. Report of the Research Results

Please submit a report describing the progress and results of the research once the study has been completed. If you are going to publish a paper based on the results of the study, please acknowledge the facilities of our institute that you have used. The sentence is as follows:

“This work was partly performed in the Cooperative Research Project Program of the Medical Institute of Bioregulation, Kyushu University.”

In addition, please submit a copy of the reprint of the paper.

If the results were obtained using the Omics Science Secure Information Analysis System (OASIS; <https://sis.bioreg.kyushu-u.ac.jp/jp/>), please include the following acknowledgment as well.

The infrastructure of Omics Science Center Secure Information Analysis System (OASIS; <https://sis.bioreg.kyushu-u.ac.jp/jp/>), Medical Institute of Bioregulation at Kyushu University provides the (part of) computational resource.

8. Research Expenses

Research expenses are allocated on the basis of screening results.

A. Collaborative research

A maximum budget of 500,000 yen, as travel and accommodation expenses, is provided. Travel expenses can be provided to research collaborators as well within the budget. In the case of instrument usage type project, research expenses are distributed as well. However, they can only cover the service fees of the institute, and cannot be allocated to personal consumable supplies and materials.

B. Research meetings

The meeting, travel, and accommodation expenses are granted.

International academic meetings: up to 1,500,000 yen

The other meetings: up to 500,000 yen

9. Handling of Intellectual Properties

Intellectual properties are handled in accordance with the Intellectual Property Handling Rules of Kyushu University.

10. Others

- (1) MIB is not responsible for any injuries or accidents during collaborative research / research meeting.
- (2) By laws and ordinances, you MUST receive an appropriate education and training or an ethical approval from either internal or external committee to meet the required ethical standards, for researches that contain genetic modifications and animal experiments, ethical issues such as human genomic information and materials, or other restricted activities.

11. Contact

Noda Tomomi /Miho Ohta (contact person)

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