

**The Guideline for Application for 2019 NIG-JOINT
(Joint Research and Research Meeting)
National Institute of Genetics,
Research Organization of Information and Systems**

1. The Guidelines for Application

(1) NIG-JOINT

National Institute of Genetics solicits Joint Research to be conducted between researchers from other universities or institutes and NIG faculties using NIG facilities and equipment on the proposed project. There are three types of research project: NIG-JOINT (A), NIG-JOINT (B) and NIG-JOINT (I).

As a general rule, travel expenses are provided to the researchers visiting NIG.

NIG-JOINT projects should be conducted during the period of time from April 1st, 2019 to March 31st, 2020. Applicants may submit applications with the same research subject as a new application up to three consecutive years.

① **NIG-JOINT (A)**

- Eligible applicants are researchers affiliated with domestic or overseas universities or research institutes. Travel expenses are provided for researchers visiting NIG for the Joint Research.
- Up to 200,000 JPY can be requested by an applicant.

② **NIG-JOINT (B)**

- Eligible applicants are researchers affiliated with domestic or overseas universities or research institutes. Travel and research expenses (supplies and other expenses used within NIG, in principle) required for the joint research are provided.
- The joint research should be planned and conducted such that applicant or his/her collaborative research members whose names are listed on the application stay at NIG for more than 7 days in total during the research period.
- Up to 1,000,000 JPY can be requested by an applicant for travel and research expenses.
- Limited numbers of applications are to be accepted compared to (A).
- As an option, you can simultaneously apply for NIG-JOINT (A), in case where (B) application is not approved. If you wish to take this option, please put a circle mark and fill in your budget for (A) (up to 200,000 JPY) in the appropriate spaces in the application.

③ **NIG-JOINT (I) – International Joint Research –**

- Eligible applicants are researchers affiliated with overseas universities or research institutes. Travel expenses are provided to researchers visiting NIG for conducting Joint Research only.
- More than 200,000JPY up to 500,000JPY can be requested by an applicant.
- Projects to be selected will be around 5.
- As an option, you can simultaneously apply for NIG-JOINT (A), in case where (I) application is not approved. If you wish to take this option, please put a circle mark and fill in your budget for (A) (up to 200,000 JPY) in the appropriate spaces in the application.

(2) Joint Research Meeting

The research meeting will be held among small number of researchers from inside and outside of NIG. The travel expenses are provided for the visit to NIG, because the meeting should be held at NIG.

The meeting must be held in the period from April 1st, 2019 to March 31st, 2020.

- Eligible applicants are NIG faculties and researchers affiliated with domestic or overseas universities or research institutes and travel expenses for attending the meeting will be provided.
- Up to 500,000JPY can be requested by an applicant.

2. Applicants

The applicants should be, as a general rule, researchers affiliated with universities, inter-university collaborative research institutes and independent administrative organizations in Japan or researchers affiliated with overseas universities or research institutes. Please note that graduate students are not eligible to apply, but they can be a member of joint researchers.

3. Application

- (1) Before submitting the application, the applicant must consult closely with the prospective NIG partner regarding project subject, title of research meeting, members, budget and other necessary matters. Research outlines and contact information of NIG researchers are described in the attached sheet.
- (2) When applying for this program, please fill out the prescribed application form and obtain approval from the head of your organization (or department). No signature or official seal is required.
- (3) Application in word format must be e-mailed to the address below. When e-mailing your application materials, please enter subject line as follows: "Application for NIG-JOINT + your name". You will be notified by e-mail within 3 working days after receiving your application documents. If you do not hear from us within a reasonable period of time, please make sure to contact us.
Prescribed form of application can be downloaded from the NIG website at:
<https://www.nig.ac.jp/nig/research-infrastructure-collaboration/nig-collaboration-grant>
【Send your application to】 kyodo-mail@nig.ac.jp (Research Promotion Team)

4. Application Deadline

Applications must arrive no later than midnight (12:00am) on December 14th, 2018.
(Japan standard Time)

5. Selection

The outcome of selection and amount of budget allocation are determined after screening by NIG committee and notified to the applicants via e-mail by the end of March 2019. Applications from those who have been adopted as research representatives, regardless of whether it is a new or a continuous project, are reviewed, taking past budget implementation status into consideration. If none of the budget has been spent in the past project, it will be a disadvantage to the applicant. If the unused budget is refunded,

accompanied by a written statement of reasons, the budget implementation status doesn't affect the selection in the next and following years.

6. Change of the Contents of Application

If there is a change in applicant's or joint research member's organization or position, or addition of a new member, it must be immediately reported to NIG representative. If applicant's affiliation is changed, a letter of approval from new organization should be submitted. (No official seal or signature is required.)

7. Expenses Provided

(1) The travel and research expenses are to be provided by NIG based on the rules of Research Organization of Information and Systems (ROIS). Because the expenses are allocated only to NIG, not to other organizations, they should be claimed through NIG representative. Accommodation fee for those who stay at our guest house will be 2,500JPY/night, and for those who stay at a hotel in the city will be 6,000JPY/night. (Outside researchers visiting NIG for attending meeting or conducting research are to stay at our guest house, as a general rule.) Please note that the travel expenses could not be fully provided in case other business or private site visits are included in the entire trip.

(2) Applicants are expected to spend the budget at an earlier stage. If you are not able to spend none of the budget due to unavoidable reason, please submit a statement of reason (free format) and refund it by the end of December. When it becomes obvious that the expenses are not used anymore, please contact Research Promotion Team at that time.

8. Submission of Research Report

The report of Joint Research or Research Meeting should be submitted within 30 days after finishing the research.

9. Publication of Research Result

You are requested to acknowledge NIG Joint Research as follows when results based on this research project are published, and send a copy of papers (PDF file is acceptable) to Research Promotion Team.

NIG-JOINT (Reference Number)

*Reference number **【Serial Number + Category (A,B or I) + year】** , will be issued in the Acceptance Notification which will be sent to the successful applicant.

Example: (Successful Applicant : Taro Iden, Reference Number : 1A2019)

“This work was supported by NIG-JOINT (1A2019) to T. Iden.”

10. Others

(1) NIG facilities and equipment are available to be used for the Joint Research and Research Meeting.

(2) When holding Research Meeting, NIG representatives are requested to post the meeting program on the NIG website and give notice to all NIG members by at least one month prior to the date of the meeting.

- (3) If gene recombination and/or animal experiments are planned, NIG representatives are requested to submit Experiment-on-Gene Recombination plan and/or Experiment-on-Animals plan application form to Research Promotion Team after acceptance of the application. As for animal experiments, researchers directly handle experimental animals are requested to apply for NIG qualification screening and undergo a training regardless of their affiliation. We strongly hope that all researchers comply with regulations and conduct the research properly.
- (4) Researchers who handle Radioisotope at NIG are requested to register as Radiation Worker before starting to handle it.
- (5) Outside researchers visiting NIG for attending meeting or conducting research are to stay at our guest house, as a general rule.
However, they can stay at a hotel in the city if the guest house is fully booked.
- (6) Ownership of intellectual property rights created in the Joint Research of NIG will be considered based on the regulations of ROIS employee invention.
- (7) NIG assures that private information for this application should be used only for screening the proposal. Regarding the accepted proposal, the representative of the research, his/her institute and the research project title will be posted on NIG website and annual reports.
- (8) As a general rule, NIG would not prepare the form of “business-trip request” for the Joint Research and Research Meeting to simplify the procedures.
We sincerely ask outside researchers visiting NIG to follow appropriate business trip procedures according to the rules of their affiliated institute.

【Contact/Inquiry】

Department of Administration

Research Promotion Team, General Affairs and Project Section

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Research outline

(Updated: January 1st,2019)

Department	Laboratory	Faculty	Research outline
Department of Informatics	Gene- Expression Analysis	OKUBO, Kousaku / Professor	“How can we make use of data and information at our finger-tip in making smarter decisions in our own contexts?” Without solving this question, all analytical and descriptive efforts that digitalize the reality end up in vain. Our tentative answer/goal for this is to develop method to enhance “fluidity” and “utility”of medical knowledge among humans and machines.
	Genetic Informatics	KAWAMOTO, Shoko /Associate Professor	We are working on research and development of databases and information retrieval system for the national bio-resource project(NBRP).
	Genome Evolution	KUROKAWA, Ken / Professor MORI, Hiroshi / Assistant Professor	We are interested in understanding about microbial genome evolution and microbial community dynamics, and we are currently reaching out in the following two major research directions; I. Facilitate the development of an integrated database “MicrobeDB.jp”, II. Microbial community dynamics.
	Biological Networks	ARITA, Masanori / Professor KAWASHIMA, Takeshi / Assistant Professor	Network analysis of metabolic pathways based on comprehensive identification and quantification of metabolites (metabolomics); Bioinformatics related to plant secondary metabolism and lipid metabolism
	Genome Informatics	NAKAMURA, Yasukazu / Professor TANIZAWA, Yasuhiro / Assistant Professor	Intelligent information technology for structural and functional annotations of large-scale nucleotide sequences.
	Research and Development of Biological Databases	TAKAGI, Toshihisa / Professor *To be retired on March 31 st ,2019	We are researching to apply distributed database technology and/or parallel-distributed computing technology to huge life science databases including DDBJ. Studies on analyzing biological data with using supercomputer.

Department of Genomics and Evolutionary Biology	DNA Data Analysis	IKEO, Kazuho / Associate Professor	Evolutionary study of genomic structure and gene expression pattern to elucidate the evolutionary mechanism of central nervous system and sensory organs. Evolutionary genomics analysis of various species. Metagenome analysis. Developing databases and computer software for biological research.
	Population Genetics	SAITOU, Naruya / Professor JINAM, Timothy / Assistant Professor	We study evolution of genes and genomes, in particular human evolution. We also develop methods for study of genome evolution.
	Plant Genetics	SATO, Yutaka / Professor TAKAHASHI, Misuzu / Assistant Professor SUZUKI, Toshiya / Assistant Professor	The goal of our research is to understand molecular mechanisms governing early processes of plant development using a series of rice embryogenesis defective mutants. Currently we are focusing on the mechanism of regulating the cell division pattern and plasticity in cellular differentiation in rice embryo.
	Evolutionary Genetics	AKASHI, Hiroshi / Professor MATSUMOTO, Tomotaka / Assistant Professor	Mechanisms of genome evolution. Especially weak selection and biosynthetic constraints.
	Human Genetics	INOUE, Ituro / Professor NAKAOKA, Hirofumi / Assistant Professor	Medical genomic study using high-throughput sequencing data is a promising procedure to create an innovate healthcare system and open a new aspect of population genetics.
	Ecological Genetics	KITANO, Jun / Professor ISHIKAWA, Asano / Assistant Professor	We use threespine stickleback fishes to investigate the genetic and molecular mechanisms underlying adaptation and speciation.
	Comparative Genomics	TOYODA, Atsushi / Project Professor	We have been conducting advanced genomics research on the plasticity of genome structure and functions using most advanced genome technology such as New-Generation Sequencers.
Department of Gene Function and Phenomics	Gene Network	SUZUKI, Emiko / Associate Professor	Combinations of molecular genetics of Drosophila and high-resolution light and electron microscopy are employed to study functional implication of structural and molecular organization of cells.

Symbiosis and Cell Evolution	MIYAGISHIMA, Shin-ya / Professor FUJIWARA, Takayuki / Assistant Professor	In order to understand endosymbiotic evolution of eukaryotes, we are studying coordinating mechanisms of eukaryotic cell and organelle/endosymbiont proliferation using algae, plants, and protists.
Model Fish Genetics	SAKAI, Noriyoshi / Associate Professor KAWASAKI, Toshihiro / Assistant Professor	We establish reliable protocols for genetically modification of zebrafish using sperm, and analyze the molecular mechanisms of spermatogenesis and early development in zebrafish.
Plant Cytogenetics	NONOMURA, Ken-ichi / Associate Professor TSUDA, Katsutoshi / Assistant Professor	We aim to elucidate the regulatory system of plant germ-cell development and chromosome kinetics, mainly using seed-sterile rice mutants.
Mammalian Neural Circuits	IWASATO, Takuji / Professor NAKAGAWA, Naoki / Assistant Professor	We are studying molecular and cellular mechanisms of neuronal circuit development in the mammals, using mouse genetics and other related methods.
Multicellular Organization	SAWA, Hitoshi / Professor	We are studying the mechanisms that produce a variety of cell types through asymmetric cell divisions using the nematode <i>C.elegans</i> .
Brain Function	HIRATA, Tatsumi / Professor KAWASAKI, Takahiko / Assistant Professor YAN, Zhu / Assistant Professor	Development of the vertebrate nervous system with special focus on neuronal network formation.
Molecular and Developmental Biology	KAWAKAMI, Koichi / Professor ASAKAWA, Kazuhide / Assistant Professor MUTO, Akira / Assistant Professor	Genetic studies on development, morphogenesis and behaviors by using a model vertebrate zebrafish.
Mammalian Development	SAGA, Yumiko / Professor KATO, Yuzuru / Assistant Professor AJIMA, Rieko / Assistant Professor	We study the early developmental events and the regulatory mechanisms during mouse embryogenesis through generation and analyses of gene-knockout and transgenic mice. We are especially interested in the organs derived from mesoderm (heart, lung, somite), and the germ cell system.
Microbial Physiology	NIKI, Hironori / Professor AOKI, Keita /Assistant Professor	We investigate higher order structure of chromosomes and their dynamics in yeast and bacteria through genetic and cell biological analysis.

	Mammalian Genetics	SHIROISHI, Toshihiko / Professor *To be retired on March 31 st ,2019 TAKADA, Toyoyuki / Assistant Professor	In order to understand genetic regulation of complex traits, such as morphogenesis and energy metabolism, we are conducting genetic analyses using mouse spontaneous mutants (variants) and genetically modified mutants.
	Mouse Genomics Resource	KOIDE, Tsuyoshi / Associate Professor TAKANAMI, Keiko /Assistant Professor	For understanding genetic basis of behavioral diversity, behavioral and genetic analyses are applied on a variety of mouse resources including wild-derived strains. We are developing genome editing methods in mice for analyzing function of genes.
Department of Chromosome Science	Epigenomics	KAKUTANI, Tetsuji / Professor TARUTANI, Yoshiaki / Assistant Professor INAGAKI, Soichi / Assistant Professor	Control and function of epigenetic gene modifications in Arabidopsis.
	Genome Dynamics	MAESHIMA, Kazuhiro / Professor IDE, Satoru / Assistant Professor HIBINO, Kayo / Assistant Professor	Our research interest lies in determining how a long string of genomic DNA is three-dimensionally organized in living cells, and how the organized genome functions during cellular proliferation, differentiation, and development. We are using a novel combination of molecular cell biology and biophysics to elucidate 3D-organization and dynamics of human genome chromatin.
	Cell Architecture	KIMURA, Akatsuki / Professor TORISAWA, Takayuki / Assistant Professor	To understand the three-dimensional architecture of the cell and its dynamics, quantitative imaging and modeling approaches are employed. Specific targets of the research are size and shape of organelles, the mechanics of cytokinesis, and cytoplasmic streaming in the <i>C. elegans</i> embryo.
	Microbial Genetics	ARAKI, Hiroyuki / Professor	Genetic and biochemical approach to elucidate molecular mechanism and regulation of eukaryotic DNA replication and checkpoint control using budding yeast
	Molecular Cell Engineering	KANEMAKI, Masato / Professor NATSUME, Toyoaki / Assistant Professor	To understand DNA transactions in human cells, we generate conditional cells using the auxin-inducible degraon technology for genetic and cytological analyses. We also develop new technologies for construction of mutant human cells.

	Invertebrate Genetics	SAITO, Kuniaki / Professor KONDO, Shu / Assistant Professor MIYOSHI, Keita / Assistant Professor	We investigate molecular mechanisms of <i>Drosophila</i> gene expression and repression through biochemical and genetic techniques. Especially, we are focusing on the small RNA pathways and chromatin regulation during germ cell development.
Center for Frontier Research	Cell Dynamics and Organization	ODA, Yoshihisa / Associate Professor	To understand the mechanism underlying plant cell wall patterning, we study the dynamic behavior of cortical cytoskeletons and small GTPases in xylem cells.
	Systems Neuroscience	KUBO, Fumi / Associate Professor	We study how visual information generates goal-directed behavior. We aim to understand the neural circuit mechanisms underlying this process using a combination of genetic, optic and behavioral approaches in zebrafish.
	Chromosome Biochemistry	MURAYAMA, Yasuto / Associate Professor	We investigate molecular mechanism underlying regulation of chromosome organization and dynamics by recapitulating their biochemical reactions using purified proteins. We now especially focus on SMC complexes.
	Quantitative Mechanobiology	SHIMAMOTO, Yuta / Associate Professor	Our laboratory studies mechanisms of force-based regulation in the mitotic spindle and the cell nucleus. Using our expertise of controlled mechanical manipulation and high-resolution fluorescence imaging, the micro-mechanics of these intracellular structures, assembled in <i>Xenopus</i> egg extracts, are quantitatively analyzed.
Advanced Genomics Center		NOGUCHI, Hideki / Project Professor	Development of new algorithms for <i>de novo</i> sequence assemblies, and analytical tools for comparative genomics employing massive data produced from next generation sequencers.
Radioisotope Unit		ANDACHI, Yoshiki / Assistant Professor	We study microRNA-mediated post-transcriptional regulation in <i>C. elegans</i> using our original methods for the detection of microRNAs and target genes.