

The Guideline for Application for 2013 Collaborative Research and Research Meeting
National Institute of Genetics,
Research Organization of Information and Systems

1. The Guidelines for Application

(1) Collaborative Research

The Purpose is to promote collaborative research between NIG faculty and researchers outside of NIG.

Based on applications from the researchers, the NIG researchers collaborate with them for conducting the research on the subject of application.

The following two categories are solicited for Collaborative Research : (A) and (B).

Collaborative Research is usually conducted during the period of time from April 1, 2012 to March 31, 2013. It can be extendable up to 3 years.

① Collaborative Research (A)

- Travel expenses only are provided for conducting the Collaborative Research within the accepted budget.
- The travel expenses are, in principle, to be paid only to the researchers who visit NIG for conducting the Collaborative Research.
- A total amount for the budget is up to 200,000JPY for each application from within Japan and up to 500,000JPY for that from overseas.

② Collaborative Research (B)

- Based on an application, both travel expenses and research expenses are provided to the researchers within the accepted budget. (For research expenses, only expendable items to be used in NIG can be bought).
- Once accepted, the representative of applicants or its collaborative researchers may stay for more than 14 days at NIG.
- Travel expenses are to be paid only when the researchers visit NIG for Collaborative Research.
- For each of applications accepted, up to 1,000,000 JPY are provided for research expense including 200,000 JPY for travel expense.
- About 10 applications are usually to be accepted
- When your application is not accepted in Collaborative Research(B), the application can be considered in Collaborative Research(A), when requested. In this case, you are requested to mention it in the application form.

(2) Research Meeting

The Purpose is to promote exchange of information between NIG faculty and researchers outside of NIG.

Based on applications from the researchers, the Research Meeting can be held in collaboration with the NIG researchers.

We provide travel expenses for visiting place where the Research Meeting is held. The Research Meeting should be held with the period from April 1, 2012 to March 31, 2013.

- Based on the application, travel expenses for the Research Meeting are to be provided.
- The Research Meeting is, in principle, held in NIG. The travel expenses are to be paid only to the non-NIG researchers who visit NIG for participating the Research Meeting.
- A total amount of money for the budget should be up to 500,000JPY per an application.

2. Exceptions

There are some exceptions as below;

(1) Collaborative Research

In the Collaborative Research only when the NIG researchers need to visit a research institution where the non-NIG researchers of Collaborative Research belong to, the travel expense can be used for it (within provided travel expenses). This can be done at any time.

(2) Research Meeting

Because Research Meeting is held in NIG, in principle, travel expenses are to be paid only to the non-NIG researchers who visit NIG. However, Research Meeting can be held at the outside of NIG, (in domestic only, when necessary.)

3. Applicants

The applicant should be, in principle, a researcher belonging to a university, an inter-university collaborative research institute and independent administrative organizations in Japan or a researcher belonging to an overseas university or a research institute.

4. Application

Please send an application form by post to the administration office with the administrative approval. (Any supervisory authority of the applicant is acceptable. In the case of the overseas applicants, the approval is exempt from this requirement.)

An application form can be downloaded from the NIG website.

<http://www.nig.ac.jp/welcome/kyoudoukenkyu/annai.html>

5. Submission of the application form

Please write “Enc. Collaborative Research Application Form” in red letters on the envelope and send it by registered or certified mail. Those affiliated with research institute outside Japan can submit an application in PDF file via e-mail.

Mailing Address

Research Promotion Team, Research Promotion Section

Department of Administration

National Institute of Genetics, Research Organization of Information and Systems

Yata1111, Mishima, Shizuoka

411-8540 JAPAN

Phone: +81-55-981-6728

E-mail: kyodo-mail@nig.ac.jp

Application Deadline:

Applications must arrive no later than 12:00 pm on January, 21 th, 2013.

(Japan standard Time)

6. Notification of the Outcome of Selection

The outcome of application will be notified to the successful candidates after screening.

The acceptance list will be also posted on NIG website.

7. Expenses Provided

Expenses will be provided by NIG within the accepted budget. The travel expenses are to be provided based on the rule of Research Organization of Information and Systems (ROIS).

8. The Report of Research

The report of Collaborative Research or Research Meeting should be submitted to the Director-General of NIG within 30 days immediately after finishing the research.

Please understand that the report might be published in an annual report of NIG.

When you write papers and make presentation within the framework of this grant, you are requested to specifically mention this grant as follows:

For Japanese : 国立遺伝学研究所共同研究 (2013A*あるいはB*)

For English : NIG Collaborative Research Program (2013-A* or B*)

(* : Reference number in the acceptance list)

In the case of thesis, it or its copy may also be submitted to the Director-General.

9. Others

(1) We strongly hope that an applicant should consult with the faculty of NIG as to the following details before submitting an application form.

(I) Collaborative Research : Proposed Research Title, expected participants,
required expenses and other necessary matters.

(II) Research Group : Name of the Research Group, purpose of the research,
proposed conducting date, expected participants,
required expenses and other necessary matters.

(2) Attached please see the document regarding the guidelines of research and the faculties in charge.

If you would like to call the faculties, please dial +81-55-981-****.

(**** : extension number)

(3) NIG makes available to our facilities for the Collaborative Research and Research Group.

(4) If you experiment for gene recombination and/or animals, you are requested to submit of Experiment-on-Gene Recombination plan and/or Experiment-on-Animals plan application form through the representative of NIG after acceptance of your application. We strongly hope that you comply with regulations and conduct the research properly.

(5) If you use Radioisotope at NIG, you are requested to register for Radiation Worker after acceptance of your application.

(6) We make the researchers who visit NIG for Collaborative Research or Research Group available to our Guest house.

(7) Regarding intellectual property created in the Collaborative Research of NIG, Ownership of the right is to be considered based on the regulations of ROIS employee invention.

(8) NIG assures that private information for this application should be used only for examining 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 a publication.

(9) Please note that NIG would not prepare the form of “business-trip request” for the Collaborative Research and Research Group because of simplicity of procedures. Please contact us mentioned below if needed.

Department of Administration

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Research outline

(Update: January 1st, 2013)

Research Department	Research Division	In charge of faculty	Extension	Research outline
Molecular Genetics	Molecular Genetics	FUKAGAWA, Tatsuo /Professor HORI, Tetsuya /Assistant Professor NISHINO, Tatsuya /Assistant Professor	6792 6744 6744	Molecular genetic, cell biological, biochemical, and structure biological methods are employed to study the mechanism for chromosome segregation during cell division.
	Mutagenesis	YAMAOKA, Fumiaki /Professor ※To be retired on March 31st, 2013	6748	There is targeted maintenance of chromosomal integration through DNA damage repair, recombination etc, especially with their linkage to ubiquitin or ubiquitin-like modification of the proteins involved in the process.
	Molecular Mechanisms	SEINO, Hiroaki / Assistant Professor	6745	I am studying molecular mechanisms of cell cycle regulation in fission yeast by genetic and biochemical approaches.
Cell Genetics	Cytogenetics	KOBAYASHI, Takehiko /Professor IIDA, Tetsushi / Assistant Professor AKAMATSU, Yufuko /Assistant Professor	6881 6882	Relationship between genome instability (especially, of repetitive sequences) and cellular functions is studied.
	Microbial Genetics	ARAKI, Hiroyuki /Professor TANAKA, Seiji / Assistant Professor HIZUME Kohji /Assistant Professor	6754 6758 6757	Genetic and biochemical approach to elucidate molecular mechanism and regulation of eukaryotic DNA replication and checkpoint control using budding yeast

Developmental Genetics	Developmental Genetics	HIROMI, Yasushi /Professor	6767	Developmental genetics of organogenesis in <i>Drosophila</i> .
		ASAOKA, Miho / Assistant Professor	6811	
		HAYASHI, Takashi / Assistant Professor	6811	
		SHIMIZU, Hiroshi / Assistant Professor	6768	Our group is currently investigating the physiological mechanism of <i>Hydra</i> and other members of phylum Cnidaria and its relation to the mechanism of pattern formation e.g. regeneration and budding.
	Neurogenetics	IWASATO, Takuji /Professor	6773	We are studying molecular and cellular mechanisms of neuronal circuit development in the mouse somatosensory system (whisker-barrel system) using mouse genetics. We are also interested in roles of alpha-chimerin in brain development and function.
		MIZUNO, Hidenobu / Assistant Professor	6777	
	Molecular and Developmental Biology	KAWAKAMI, Koichi /Professor	6740	Genetic studies on development, morphogenesis and behaviors by using a model vertebrate zebrafish.
		ASAKAWA, Kazuhide / Assistant Professor	6739	
		MUTO, Akira /Assistant Professor		
Population Genetics	Population Genetics	SAITOU, Naruya /Professor	6790	We study evolution of genes and genomes, in particular human evolution. We also develop methods for study of genome evolution.
		SUMIYAMA, Kenta / Assistant Professor	6787	

	Evolutionary Genetics	AKASHI, Hiroshi /Professor OSADA, Naoki / Assistant Professor	6793 5820	Mechanisms of genome evolution. Especially weak selection and biosynthetic constraints.
Integrated Genetics	Human Genetics	INOUE Ituro /Professor HOSOMICHI, Kazuyoshi /Assistant Professor	6795 6797	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.
	Agricultural Genetics	KAKUTANI, Tetsuji /Professor TARUTANI, Yoshiaki / Assistant Professor INAGAKI, Soichi /Assistant Professor	6801 6807	Control and function of epigenetic gene modifications in Arabidopsis.
	Brain Function	HIRATA, Tatsumi /Associate Professor KAWASAKI, Takahiko / Assistant Professor	6721 6721	Development of the vertebrate nervous system with special focus on neuronal network formation.
Center for Frontier Research	Molecular Function	KANEMAKI, Masato /Associate Professor	5830	We aim to understand the mechanism of chromosome replication and the cell cycle regulation in animal cells by analyzing conditional cell lines using molecular genetic and cell biological methods. We also develop techniques for the construction of cell lines required for the studies of animal cells.

	Motor Neural Circuit	HIRATA, Hiromi /Associate Professor	5825	Genetic and physiological analysis on motor development by using a vertebrate model zebrafish. Specific aim is to understand and regulate intrinsic and acquired synaptogenesis, circuit formation and muscle development.
	Symbiosis and cell evolution	MIYAGISHIMA, Shin-ya /Project Associate Professor	9411	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
	Ecological Genetics	KITANO, Jun /Project Associate Professor /	9415	We use threespine stickleback fishes to investigate the genetic and molecular mechanisms underlying adaptation and speciation.
	Centrosome Biology	KITAGAWA, Daiju /Project Associate Professor	5828	We mainly focus on understanding the mechanisms of centrosome duplication by using the combination of innovative and multi-disciplinary approaches. We are utilizing <i>C. elegans</i> embryos and human cell culture as model systems.
Genetic Strains Research Center	Mammalian Genetics	SHIROISHI, Toshihiko / Professor	6818	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.
		TAMURA, Masaru / Assistant Professor	6816	
TAKADA, Toyoyuki / Assistant Professor	6820			
	Mammalian Development	SAGA, Yumiko / Professor	6829	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.
		KATO, Yuzuru / Assistant Professor	6815	
		AJIMA, Rieko /Assistant Professor		

Mouse Genomics Resource	KOIDE, Tsuyoshi /Associate Professor TAKAHASHI, Aki /Assistant Professor	5843	For understanding genetic basis of behavioral diversity, behavioral and genetic analyses are applied on a variety of mouse resources including wild-derived strains.
Model Fish Genomics Resource	SAKAI, Noriyoshi /Associate Professor SHINYA, Minori / Assistant Professor	5848 5849	We establish reliable protocols for genetically modification of zebrafish using sperm, and analyze the molecular mechanisms of spermatogenesis and early development in zebrafish.
Plant Genetics	KURATA, Nori / Professor KUBO, Takahiko / Assistant Professor	6808 6802	We perform analyses of genetic programs of reproductive and embryonic developmental process, as well as studies on the mechanism of reproductive isolation in rice. Wild species resources of rice are also used for evolutionary and diversity studies.
Microbial Genetics	NIKI, Hironori / Professor AOKI, Keita /Assistant Professor	6870	We investigate higher order structure of chromosomes and their dynamics in yeast and bacteria through genetic and cell biological analysis.
Invertebrate Genetics	UEDA, Ryu / Professor KONDO, Syu /Assistant Professor	6823	Genome-wide RNAi mutant fly library is established to study genome function in a variety of biological traits of fly development.
Genetic Informatics	YAMAZAKI, Yukiko /Associate Professor	6885	As the information center of the genetic resources, we have been constructing databases and continuously inventing better way to distribute data in order to utilize the resources to its fullest potential.
Genome Biology	KOHARA, Yuji / Professor ANDACHI, Yoshiki / Assistant Professor	6854 6860	We are performing a systematic analysis of expression and function of the genome of the nematode C.elegans, aiming at understanding of the gene network for development.

Structural Biology Center	Biological Macromolecules	MAESHIMA, Kazuhiro / Professor HIRATANI Ichiro /Assistant Professor	6864	Our research interest lies in determining how a long string of genomic DNA is three-dimensionally organized in mitotic chromosomes and the nucleus, 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 /Associate Professor KIMURA, Kenji /Assistant Professor	5854	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 <i>C. elegans</i> embryo.
	Multicellular Organization	SAWA, Hitoshi /Professor IHARA, Sinji /Assistant Professor	6845	We are studying the mechanisms that produce a variety of cell types through asymmetric cell divisions using the nematode <i>C.elegans</i> .
	Biomolecular Structure	SHIRAKIHARA, Yasuo /Associate Professor ITO,Hiroshi / Assistant Professor	6887 6862	We determine the three dimensional atomic structure of proteins, nucleic acids or their complexes by x-ray diffraction analysis in order to understand the working mechanism of the targets.

	Gene Network	SUZUKI,Emiko /Associate Professor	6812 6813	Combinations of molecular genetics of <i>Drosophila</i> and high-resolution light and electron microscopy are employed to study functional implication of structural and molecular organization of neuronal cells, with particular focus on neuronal network formation.
Center for Information Biology	DNA Data Analysis	GOJOBORI, Takashi /Professor IKEO,Kazuho /Associate Professor NOZAWA, Masafumi /Assistant Professor	6847 6851 6852	Evolutionary study of genomic structure and gene expression pattern of animals to elucidate the evolutionary mechanism of central nervous system and sensory organs. Evolutionary genomics analysis of various species such as <i>Drosophila</i> and viruses. Developing databases and computer software for biological research.
	Gene Function Research			
	Genome Informatics	NAKAMURA, Yasukazu /Professor KAMINUMA,Eli / Assistant Professor	6859 6859	Intelligent information technology for structural and functional annotations of large-scale nucleotide sequences.
	Research and Development of Biological Databases	TAKAGI, Toshihisa / Professor	5821	We are researching to apply distributed database software technology and/or parallel-distributed computing software technology to huge Life Science Databases such as DDBJ.
	Gene- Expression Analysis	OKUBO,Kousaku / Professor OGASAWARA, Osamu /Assistant Professor	5838 9450	Representation of Bio Medical knowledge Analysis of gene expression data and construction of integrated databases, construction of a database of data analysis methods, and construction of theoretical models of gene expression evolution

	Comparative Genomics	FUJIYAMA, Asao /Professor TOYODA, Atsushi /Project Associate Professor	6788	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.
Experimental Farm		NONOMURA, Ken-ichi / Associate Professor MIYAZAKI, Saori /Assistant Professor	6872 6874	We aim to elucidate the regulatory system of plant germ-cell development and chromosome kinetics, mainly using seed-sterile rice mutants.
Center for Advanced Genomics		NOGUCHI, Hideki /Project Associate Professor	9459	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.

Research Department	Research Division	In charge of faculty	Extension	Research outline
Adjunct Faculty	Nucleic Acid Chemistry	EARNSHAW, William C. /Professor	6870	Studies of mitotic chromosome structure and function.
		MARKO, John F. / Professor	6748	Physics of large-scale DNA organization.
	Cytoplasmic Genetics	BOCCARD, Frédéric / Professor	6870	Dynamics of Bacterial Chromosome
		UEDA, Hiroki / Professor	6754	Systems Biology of "Time"
	Physiological Genetics	STERN, David L. / Professor	6767	Genetic causes of evolution of morphology and behavior
		FURLONG Eileen E.M / Professor	6767	Transcriptional control during development.
	Theoretical Genetics	von HAESELER, Arndt /Professor	6790	Evolutionary Bioinformatics.
		CLARK, Andrew G. /Professor	6793	Genetic basis of adaptive variation in natural populations
	Applied Genetics	MARTIENSSEN Robert A. / Professor	6801	Inheritance and reprogramming of heterochromatin with small RNA.
		TSUJI, Shoji / Professor	6788	Elucidation of the mechanisms of brain diseases based on personal genome analysis.