

# EDGE Webinar

## Enabling Discovery through Genomics

Presenters:

Patrick Abbot (NSF/IOS)  
Ford Ballantyne (NSF/DEB)  
Steve Ellis (NSF/DBI)  
Anthony Garza (NSF/MCB)  
Ted Morgan (NSF/IOS)  
Diane Okamuro (NSF/IOS)  
Floh Thiels (NSF/IOS)  
Jennifer Troyer (NIH/NHGRI)

**15 January 2021**

[BIOEDGE@NSF.GOV](mailto:BIOEDGE@NSF.GOV)



# EDGE Webinar Outline

## 1. Program Overview

- a. Purpose and goals
- b. Examples of programmatic fit
- c. Partnership with the National Human Genome Research Institute

## 2. Submission Requirements

- a. Deadline
- b. Proposal sections

## 3. Review Criteria

- a. NSF merit review criteria
- b. Solicitation-specific criteria
- c. NIH review criteria

## 4. Questions



# Program Overview

## Purpose

Enable advancement of understanding the relation between genomes and phenomes—relevant to *Understanding the Rules of Life: Predicting Phenotype*

## Goals

To support:

- (1) development of **tools**, approaches, and infrastructure for testing cause and effect hypotheses between gene function and phenotypes in **organisms for which such methods are presently unavailable**
- (2) hypothesis-driven **research** that tests cause-and-effect relations between genotype(s) and phenotypes **in diverse organisms within the context (environmental, developmental, social, and/or genomic) in which they function**. Use of traditional model organisms is permitted, but proposals must **demonstrate the generalizability of the results** beyond the focal species across contexts.



# Program Overview

## Functional Genomics Tools (FGT) track

For example:

- Development of mutant libraries and/or high-quality reference genomes
- Generalizable high-throughput phenotyping methods
- Innovative approaches for manipulating individual genes or multiple genes simultaneously
- Innovative approaches to test gene function in targeted, single cells in organisms
- Innovative approaches for establishing function of single or networks of genes

## Complex Multigenic Traits (CMT) track

For example:

- Systems-level analysis of the gene regulatory networks underlying complex traits
- Innovative analytical approaches to linking genes and complex traits
- Elucidation of the causal connections across levels of biological organization that underlie complex multigenic traits
- Elucidation of multi-genome/epigenome interactions with the environment, with the goal of predicting complex organismal phenotypes across contexts

**Use of Diverse Organisms to Advance Functional Genomics**



# Program Overview

## Both tracks:

Higher priority will be given to projects that:

- **cross disciplines** within biology, combining organismal biology with cellular, evolutionary, or ecological research
- demonstrate approaches that are **generalizable** beyond single-gene, single-trait, or single-diseases relevant phenotypes

Program **considers applications** that:

- can demonstrate utility or relevance to human or disease-relevant model organisms
- will advance the science of genomic medicine
- will incorporate genomics to improve the effectiveness of healthcare

**Use of Diverse Organisms to Advance Functional Genomics**



# Program Overview

## Functional genomics EDGE does not support:

- Genome-scale research or tool development for agriculturally relevant plant species
- Proposals focused exclusively on bioinformatic tool development

**Questions about your project's fit for the EDGE Program  
and for which track?**

Send a **brief synopsis** (~ 1 page) to [BIOEDGE@nsf.gov](mailto:BIOEDGE@nsf.gov)



# Program Overview

## NSF program in partnership with the National Human Genome Research Institute of the National Institutes of Health

To:

- Promote scientific exchange across communities traditionally supported by the respective agencies and allow for synergies that will accelerate advancement in functional genomics
- Leverage evolutionary relationships in advancing understanding of the functional mechanisms that connect genotype-to-phenotypes and the ability to predict phenotype
- Strengthen the links between fundamental research and societal needs



# Program Overview

## National Human Genome Research Institute



Supports development of resources, approaches, and technologies that will accelerate genomic research

- Structure of genomes, biology of genomes, genomics and biology of disease
- Use of genomics to advance the science of medicine and improve the effectiveness of healthcare
- Genomic research in the ethical, legal and societal implications of genomics and genetics research, bioinformatics, technology development, and research training and career development
- Generalizable methods and knowledge about genomics in relation to human health

NHGRI will consider EDGE applications that can demonstrate ***utility or relevance to human health or disease***, provided they also focus on the ***development of methods or novel applications*** that demonstrate ***approaches that are generalizable*** beyond single-gene, single-trait, or single-disease relevant phenotypes



# Submission Requirements

(Section V of the solicitation)

## Proposal submission deadline

**16 March 2021, 5 PM submitter's local time**

## Title

Functional Genomic Tools track: "FGT:...."

Complex Multigenic Traits track: "CMT:...."

## Titled sub-sections of the Project Description

FGT and CMT proposals:

- Intellectual Merit
- Broader Impacts

additional sub-section for FGT proposals:

- Research Community Impact



# Submission Requirements

(Section V of the solicitation)

## Supplementary Documents

FGT or CMT proposals that involve multiple organizations:

- Project Management Plan

Additional Supplementary Document for FGT proposals:

- Dissemination and Education Plan

Where relevant:

- Human Subjects Protection
- Vertebrate Animal Section

## Data Management Plan

required per the Proposal and Award Policies and Procedures Guide NSF 20-1



# Submission Requirements

(Section V of the solicitation)

## Letters of Collaboration

- prescribed format per the Proposal and Award Policies and Procedures Guide NSF 20-1
- **letters of support are not allowed**

## Budget and Duration

- up to \$2 million in combined direct costs (summed over all components of the project if the proposal is a collaborative submission)
- up to 4 years



# Review Criteria

(Section VI of the solicitation)

**Intellectual Merit** – Potential to advance knowledge

**Broader Impacts** – Potential to benefit society and contribute to the achievement of specific societal outcomes

Elements considered in the review for both criteria:

- What is the potential for the proposed activity to
  - Advance knowledge and understanding within its own field or across different fields (**Intellectual Merit**); and
  - Benefit society or advance desired societal outcomes (**Broader Impacts**)?
- To what extent do the proposed activities suggest and explore creative, original, or potentially transformative concepts?
- Is the plan for carrying out the proposed activities well-reasoned, well-organized, and based on a sound rationale? Does the plan incorporate a mechanism to assess success?
- How well qualified is the individual, team, or organization to conduct the proposed activities?
- Are there adequate resources available to the PI (either at the home organization or through collaborations) to carry out the proposed activities?



# Review Criteria

(Section VI of the solicitation)

## Solicitation-specific review criteria

For **FGT** proposals:

- Potential catalytic impact on advancing research and research communities using organisms named in the proposal
- Potential catalytic impact of the enabling tools, approaches, and infrastructure on advancing research and research communities
- Quality and potential for rapid and high impact of the Dissemination and Education Plan

For **CMT** proposals:

- Extent to which the proposed hypotheses test or demonstrate causal links between genomes and phenomes
- Potential to advance theory and understanding of complex multigenic trait expression
- Potential to inform predictive understanding of complex phenotypes based on genomic information
- Degree to which generalizability of the conclusions across diverse research organisms will be demonstrated



# Review Criteria

(Section VI of the solicitation)

## Solicitation-specific review criteria

For FGT or CMT proposals that involve **multiple organizations**:

- Quality of the Project Management Plan and likelihood of successful project coordination

### NIH Review Criteria:

- Overall impact
- Significance
- Investigator(s)
- Innovation
- Approach
- Environment

If applicable:

- Protection of human subjects
- Vertebrate animals
- Biohazards
- Budget and period of support



# Process at-a-glance

1. Proposal submission in accordance with submission requirements specified in Solicitation NSF 21-546
2. Proposal review managed by NSF in consultation with NHGRI
3. Identification of meritorious proposals that may be recommended for funding by either NSF or NIH
  - a. Proposals selected for funding by NSF will be handled in accordance with standard NSF procedures
  - b. Proposals selected for funding consideration by NIH will be invited to submit reformatted applications to NIH's Center for Scientific Review

**Proposal submission deadline**  
**16 March 2021, 5 PM submitter's local time**



# EDGE Webinar

EDGE Program page:

[https://nsf.gov/funding/pgm\\_summ.jsp?pims\\_id=505480](https://nsf.gov/funding/pgm_summ.jsp?pims_id=505480)

EDGE contact: [BIOEDGE@NSF.GOV](mailto:BIOEDGE@NSF.GOV)

