

NIST Genome Editing Consortium Overview

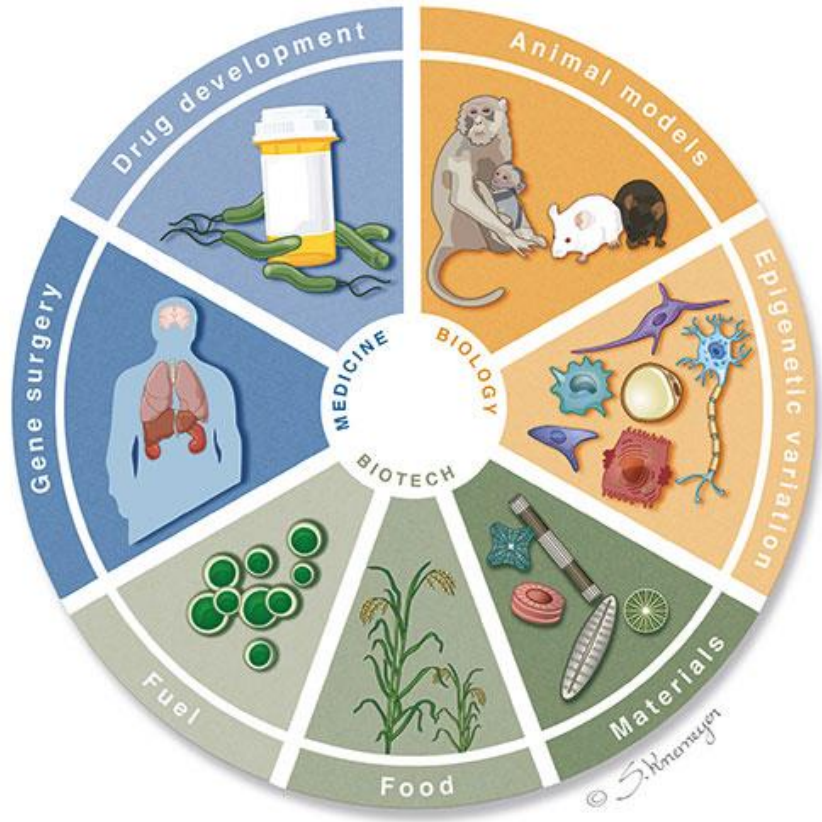
Samantha Maragh
Leader, Genome Editing Program
Samantha@nist.gov

NIST
National Institute of
Standards and Technology
U.S. Department of Commerce

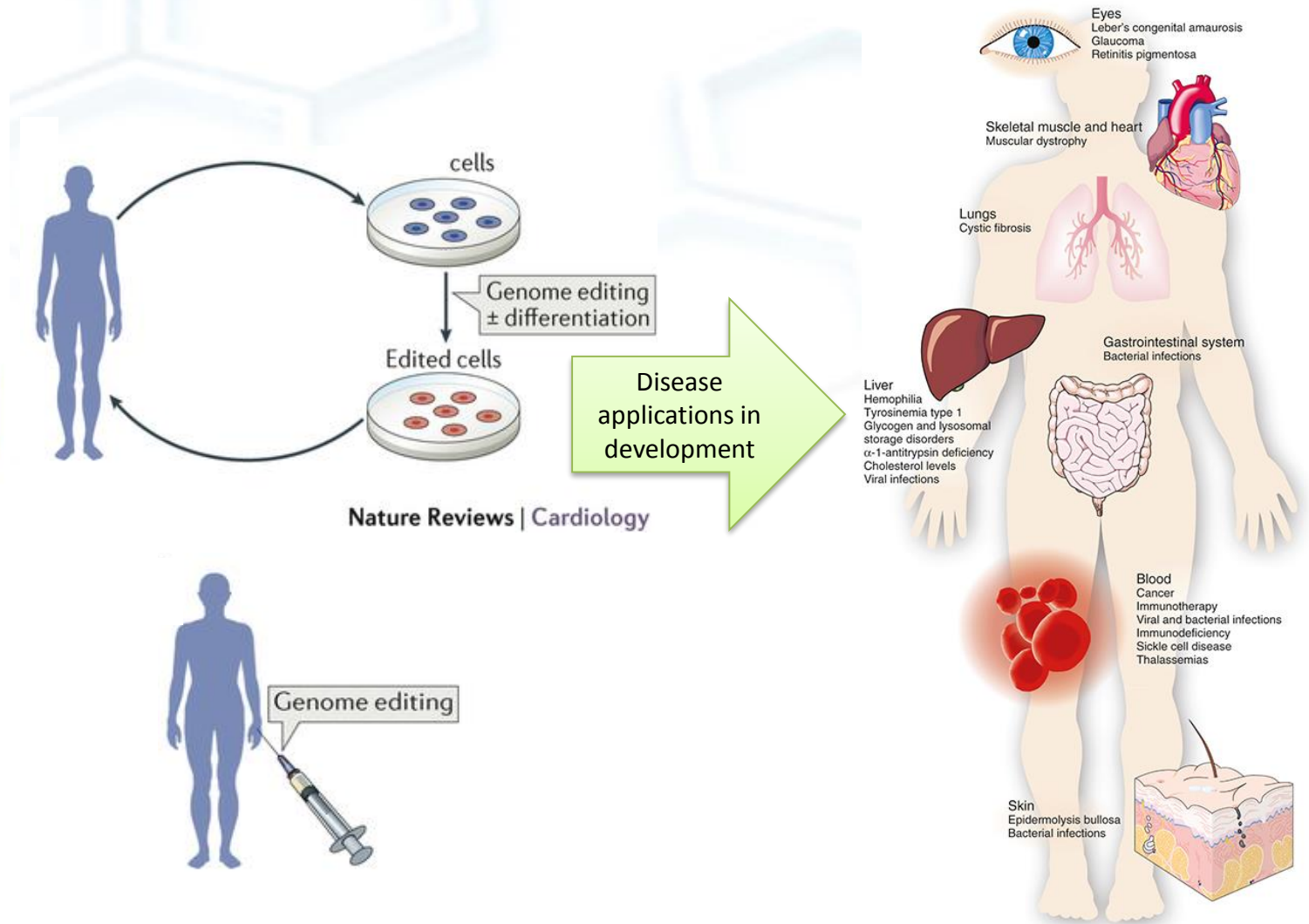
**MATERIAL
MEASUREMENT
LABORATORY**

Building Confidence in Biological Measurements

How did we get here?



REGENERATIVE MEDICINE/CELL/GENE THERAPY EXAMPLE



How did we get here?



Sponsors

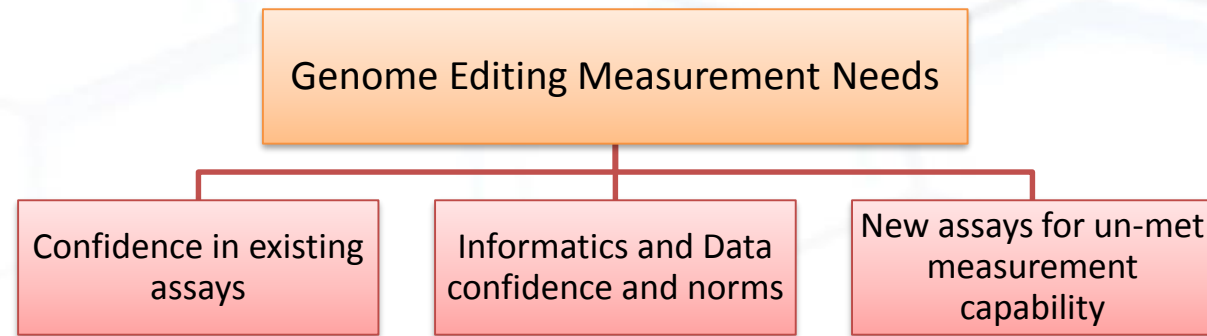


Genome Editing Standards Workshop
May 2, 2016 NIST- hosted

Where is confidence needed?

What is needed to achieve this confidence in this context?

65 participants at the 2016 NIST Genome Editing Standards Workshop identified pre-competitive standards and measurements needed to establish greater confidence in characterization of genome editing outputs.



Leading recurring needs shared from the community

- How to quantify and reliably detect on-target and off-target genome editing?
- How to compare existing off-target location discovery assays?
- How to determine reliability of validating sequence variants?
- How to compare different genome editing reagents?
- Agreement on data reporting and informatics data formats
- A standard data set to test and understand comparability of informatics pipelines

How did we get here?

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NIST Genome Editing Measurements and Standards Discussion
at the Keystone Precision Genome Engineering Conference
Jan 11th 2017

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Focus group of experts supported a path forward of a NIST led Genome Editing Consortium with 3 focus areas:

1. Specificity Measurements
2. Data & Metadata
3. Lexicon

Some of the participants in the process:



NIST Genome Editing Consortium (now accepting members)

MISSION

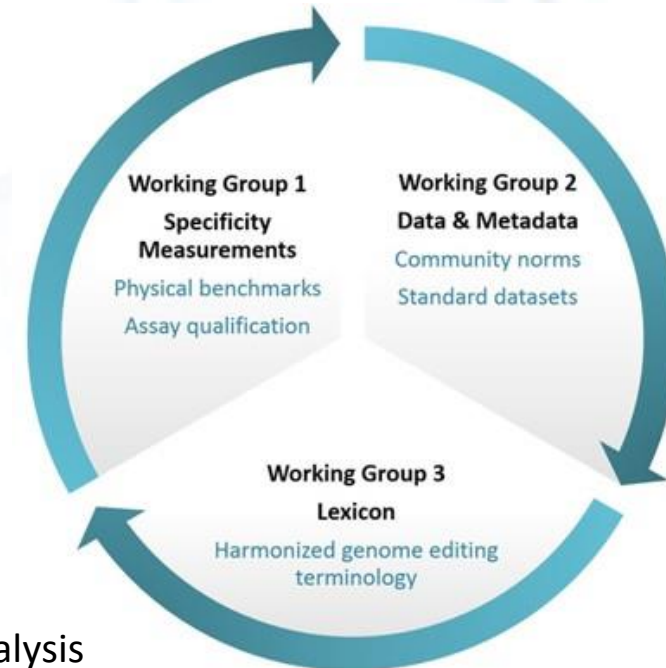
Convene experts across academia, industry, non-profit & government to address the measurements and standards needed to increase confidence and lower the risk of utilizing genome editing technologies in research and commercial products.

CONSORTIUM GOALS

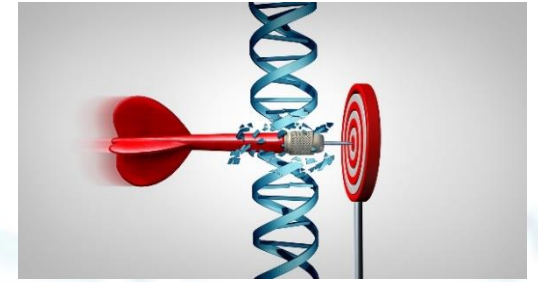
- Qualify genomic assays used to evaluate genome editing outputs
- Develop reference materials
- Generate reference data and standard data formats
- Develop suggested minimal information reporting for public studies
- Generate a common lexicon for genome editing studies

CONSORTIUM IMPACT

- Shared genome editing relevant reference materials
- Community norms for minimum data reporting and tools for supporting data analysis
- Improved understanding of and confidence in detecting and quantifying on- and off-target genome editing
- Harmonized genome editing lexicon



NIST Genome Editing Consortium Working Groups



Working Group 1: Specificity Measurements

- NIST qualification of existing genomic assays used to identify on-target and off-target genome editing genomic locations and sequence variants
- Develop & qualify reference materials

Working Group 2: Data and Metadata

- Determine community norms for minimum information reporting
- Identify community norms for data formats and tools for benchmarking data analysis (e.g. in silico data sets, experimental data sets)

Working Group 3: Lexicon

- Identify terms and related definitions to form a common genome editing community lexicon

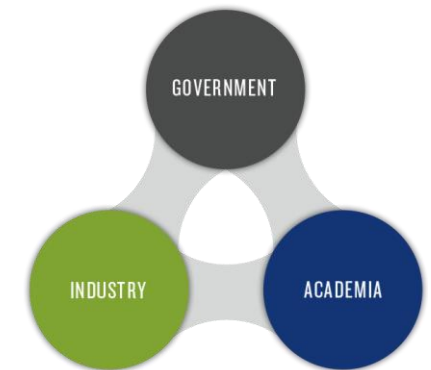
NIST Genome Editing Consortium (now accepting members)

WHY NIST?

- Cross-disciplinary expertise in engineering, and the physical, information, chemical, and biological sciences
- As a non-regulatory agency of the U.S. Department of Commerce, NIST does not impose standards; standards are accepted by consensus
- Neutral convener for industry consortia, standards development organizations, federal labs, universities, public workshops, and interlaboratory comparability testing

MODEL

- Convenes industry, academia, and government to identify and address measurement and standards needs across the genome editing field
- Enables members to work with NIST to develop measurement solutions and standards
- Leverages NIST expertise in measurement science, standards development, reference materials, technology development, and basic research
- Collaborates with related programs at other federal agencies



NIST Genome Editing Consortium

*MEMBER BENEFITS

- Access to a neutral forum for addressing pre-competitive needs
- Participation in the development of experimental benchmarks, guidelines and terminology
- Access to tools developed by the consortium ahead of public release
- Institutional representation on consortium steering committee

Membership Process:

- complete a [Letter of Interest](#).
- NIST will reach out to interested parties to initiate signing a Cooperative Research and Development Agreement (CRADA). Consortium CRADA is identical for all participants.
- *Membership fee of \$20K or in-kind support of equivalent value to join. These contributions will supplement the NIST resources to enable us to generate and qualify the pre-competitive standards, norms and data the consortium members determine are needed to push the field forward and support innovation.

website: <https://go.usa.gov/xnV3> | email: samantha@nist.gov (Samantha Maragh)

NIST Genome Editing Consortium + public workshops

At these workshops, consortium members will:

- Discuss the progress of the consortium work
- Continue to get broad input to update or refine the consortium work plan
- Continue to broadly solicit consortium membership
- Invite members to participate in work plan implementation, planned future experiments, and analysis

NIST-FDA Genome Editing Workshop April 23-24, 2018

Purpose: convene key stakeholders from industry, academia and government, to define a path forward for developing pre-competitive solutions to address the wet lab, dry lab and documentary benchmarks, norms and standards needed for genome editing- particularly the needs to support development of gene therapies.

Format: general session talks with discussion + breakout sessions to get detailed feedback and engagement.

Goals:

1. clarify regulatory perspectives
2. Inform on the path forward for the NIST Genome Editing consortium-
3. Get public input on path forward
 - consortium members working with NIST will determine how to proceed and execute to develop the needed products.

NIST-FDA Genome Editing Workshop April 23-24, 2018

Agenda Overview

DAY 1

Morning session 1: NIST, FDA, NIH SCGE

Break

Morning session 2: Expert perspectives & panel discussion

Lunch

Afternoon session 1: Plan for path forward presentations

Break

Afternoon session 2: Breakout sessions & end of Day 1

DAY 2

Morning session 1: Technology Talks (approaches for DNA off-target and genomic rearrangement detection)

Break

Morning session 2: Reports from breakout sessions, next steps, and wrap-up

Adjourn