

Future Opportunities for Genome Sequencing and Beyond: A Planning Workshop for the National Human Genome Research Institute

July 28-29, 2014

Bethesda North Marriott Hotel and Conference Center Bethesda, Maryland

AGENDA

Objectives:

- 1. Discuss the scientific questions and opportunities that can be substantially addressed by large-scale genomics studies, starting with genome sequencing but also considering other genomic technologies.
- 2. Consider options for future NHGRI programs that would address these questions and opportunities.

Monday, July 28, 2014

8:30 a.m. - 9:00 a.m. Welcome and Charge for the Workshop Salons A & B

Eric Green, M.D., Ph.D.

National Human Genome Research Institute, NIH

9:00 a.m. - 9:15 a.m. Background and Orientation

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

Part 1: The Science of the Current NHGRI Genome Sequencing Program

9:15 a.m. - 9:45 a.m. Discovering Variants Conferring Risk for Common Diseases

Michael Boehnke, Ph.D. University of Michigan

9:50 a.m. - 10:20 a.m. Discovering the Genomic Bases of Mendelian Diseases

Roderick McInnes, M.D., Ph.D.

Lady Davis Institute for Medical Research

10:25 a.m. - 10:55 a.m. Genome Sequencing for Clinical Care

Dan Roden, M.D. Vanderbilt University

11:00 a.m. - 11:20 a.m. **Break** (on your own)

11:20 a.m. - 11:50 a.m. Functional Genomics at Scale

Joseph Ecker, Ph.D.

Howard Hughes Medical Institute and Salk Institute for Biological Studies

11:50 a.m. - 12 noon Setting the Stage for the Discussion

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

12 noon - 1:15 p.m. Lunch (on your own)

1:15 p.m. - 2:15 p.m. Discussion and Refinement of Opportunities

Ewan Birney, Ph.D.

European Molecular Biology Laboratory

Part 2: How Should Scale Evolve?

2:15 p.m. - 2:45 p.m. Introduction to Breakout Groups:

What Are the "Big Challenges" That NHGRI Should Pursue

in the Next ~5 Years? William M. Gelbart, Ph.D. Harvard University

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

3:00 p.m. - 5:00 p.m. Breakout Groups (2 hours each)

Group 1: Understanding the Genetic Architecture of Health

and Disease at Scale Eric Boerwinkle, Ph.D.

The University of Texas Health Science Center at Houston

Mike Bamshad, M.D. University of Washington

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

Group 2: Integrating Genomic Variant Discovery with Function

Richard Myers, Ph.D.

HudsonAlpha Institute for Biotechnology

Mark Gerstein, Ph.D. Yale University

Elise Feingold, Ph.D.

National Human Genome Research Institute, NIH

Mike Pazin, Ph.D.

National Human Genome Research Institute, NIH

Linden Oak

Brookside A

Salons A & B

	Group 3: Clinical Genome Sequencing at Scale Heidi Rehm, Ph.D. Partners Healthcare	Forest Glen
	Sharon Emma Plon, M.D., Ph.D. Baylor College of Medicine	
	Lucia Hindorff, Ph.D., M.P.H. National Human Genome Research Institute, NIH	
	Carolyn M. Hutter, Ph.D. National Human Genome Research Institute, NIH	
	Group 4: Comparative and Evolutionary Genomics Andrew Clark, Ph.D. Cornell University	Brookside B
	Evan E. Eichler, Ph.D. University of Washington	
	Michael W. Smith, Ph.D. National Human Genome Research Institute, NIH	
5:15 p.m 5:30 p.m.	Reconvene: Issues Arising, Logistics, Evening Session Adam Felsenfeld, Ph.D. National Human Genome Research Institute, NIH	Salons A & B
5:30 p.m 7:30 p.m.	Dinner (on your own)	
7:30 p.m 9:00 p.m.	Challenge Talks (10-minute talk/5-minute question)	Salons A & B
7:30 p.m 7:45 p.m.	Peptide Display and T-Cell Recognition Project David Haussler, Ph.D. University of California, Santa Cruz	
7:45 p.m 8:00 p.m.	Using the Negative Correlation Between Polygenic and Rare Variant Burdens for Common Disease To Improve Study Efficiency and Power Nancy Cox, Ph.D. The University of Chicago	
8:00 p.m 8:15 p.m.	The Human Cell Atlas Aviv Regev, Ph.D. Broad Institute	
8:15 p.m 8:30 p.m.	Measuring the Functional Consequences of Very Large Numbers of Human Genetic Variants Jay Shendure, M.D., Ph.D. University of Washington	

8:30 p.m. - 8:45 p.m. Not Your Father's PDF: New Forms of Knowledge Representation for

Genomic Sequence Daniel Masys, M.D. University of Washington

9:00 p.m. Breakout Group Co-Chairs Meet To Coordinate Summaries

Breakout Co-Chairs and Staff Only

Tuesday, July 29, 2014

Part 3: Moving From Ideas to Implementation

7:30 a.m. - 8:30 a.m. Staff Meets with "Scenarios" Co-Chairs To Develop Discussions

(Breakout Co-Chair Coordination if Needed)

8:30 a.m. - 10:10 a.m. Breakout Reports From Day 1 (15-minute report/10-minute discussion)

Salons A & B

Salons A & B

Breakout Group Co-Chairs

8:30 a.m. - 8:55 a.m. Group 1: Understanding the Genetic Architecture of Health

and Disease at Scale

8:55 a.m. - 9:20 a.m. Group 2: Integrating Genomic Variant Discovery with Function

9:20 a.m. - 9:45 a.m. Group 3: Clinical Genome Sequencing at Scale

9:45 a.m. - 10:10 a.m. Group 4: Comparative and Evolutionary Genomics

10:15 a.m. - 10:30 a.m. **Break** (on your own)

10:30 a.m. - 10:45 a.m. Introduction: "Scenarios"

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

William M. Gelbart, Ph.D. Harvard University

10:45 a.m. - 1:30 p.m. Alternate "Scenarios" (5-minute introduction/25-minute discussion)

10:45 a.m. - 11:15 a.m. Element 1: Understanding the Genetic Architecture of Health

and Disease at Scale Len Pennacchio, Ph.D.

Lawrence Berkeley National Laboratory

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

11:15 a.m. - 11:45 a.m. Element 2: Integrating Genomic Variant Discovery with Function

Carlos Bustamante, Ph.D.

Stanford University

Mike Pazin, Ph.D.

National Human Genome Research Institute, NIH

Elise Feingold, Ph.D.

National Human Genome Research Institute, NIH

12 noon - 1:00 p.m. Lunch (on your own)

1:00 p.m. - 1:30 p.m. Element 3: Clinical Genome Sequencing at Scale

James Philip Evans, M.D., Ph.D.

The University of North Carolina at Chapel Hill

Lucia Hindorff, Ph.D., M.P.H.

National Human Genome Research Institute

1:30 p.m. - 2:30 p.m. Discussion: Combining Strategic and Tactical Considerations

Rex Chisholm, Ph.D., M.S. Northwestern University

Adam Felsenfeld, Ph.D.

National Human Genome Research Institute, NIH

Jeffery A. Schloss, Ph.D.

National Human Genome Research Institute, NIH

2:30 p.m. - 3:00 p.m. Wrap-up

Eric Green, M.D., Ph.D.

National Human Genome Research Institute, NIH

3:00 p.m. Adjournment