NIH Policy for Data Sharing of NIH Supported Genome Wide Association Studies

Elizabeth G. Nabel, M.D.

Director, National Heart, Lung, and Blood Institute

Council of Public Representatives

April 18, 2008





NIH GWAS Policy

DEPARTMENT OF HEALTH AND HUMAN SERVICES

National Institutes of Health

Policy for Sharing of Data Obtained in NIH Supported or Conducted Genome-Wide Association Studies (GWAS)

AGENCY: National Institutes of Health,

ACTION: Notice.

Background

The NIH is interested in advancing genome-wide association studies (GWAS) to identify common genetic factors that influence health and disease. For the purposes of this policy, a genome-wide association study is defined as any study of genetic variation across the entire human genome that is designed to identify genetic associations with observable traits (such as blood

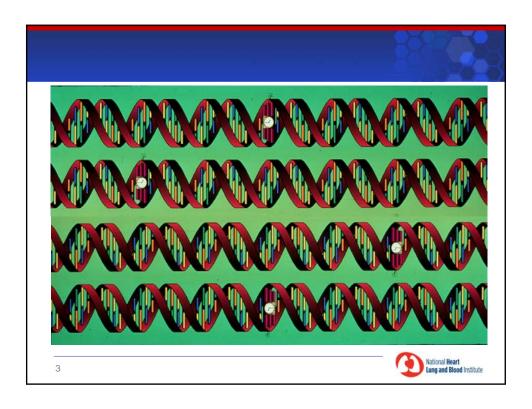
Federal Register: August 28, 2007 Policy Effective: January 25, 2008

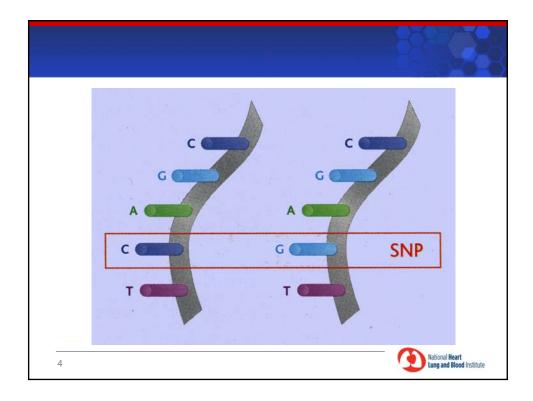
Us Department of Health & Human Services Chica of Extracruzal Research Extracruzal Research Extracruzal Research Roma About Creatis Founding Opportunities Frunding Opportunities Fru

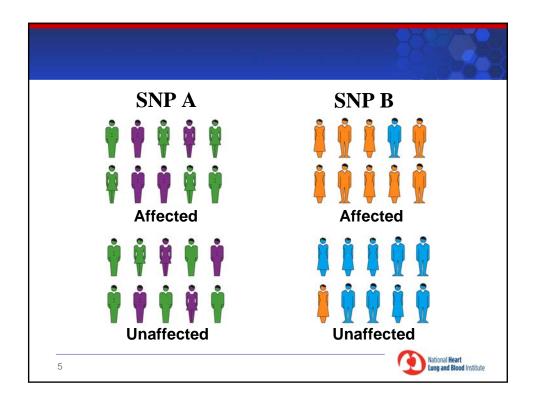
NIH OER Website:

http://grants.nih.gov/grants/gwas/index.htm





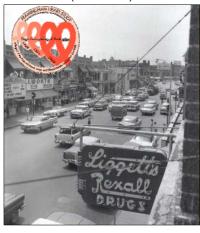






Framingham Heart Study

Downtown Framingham, MA (circa 1960)



Risk Factors for Heart Attack and Stroke

- High blood pressure
- High cholesterol
- Cigarette smoking
- Diabetes mellitus
- Parental or sibling history
- Obesity

7

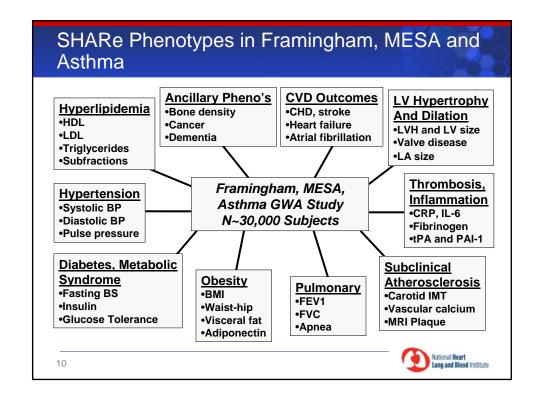


Framingham Heart Study: Population-Based Family Study 1948 + 1958 **→1968** →1978 →1988 -→ 1998 2008 948 Original cohort: n=5209 men and women (ages 28-62) 1644 spouse pairs, 596 extended families 2008 1972 Offspring study: n=5124 men and women (ages 5-70) 1576 spouse pairs, 3514 biological offspring Third Generation study: 2002 n=3500 men and women

Framingham SNP Health Association Resource (SHARe)

- Genotypes: ~10,000 Caucasians from 3 generations - Affymetrix 500,000 SNP chip
- Phenotypes: >1,000 risk factor, subclinical and clinical CV phenotypes, from 60 years of exams
- Genotypes and phenotypes placed in a webbased dataset, called dbGaP, maintained at the NIH
- Framingham SHARe dataset contains 5.5 billion genotypes, >5.5 trillion association tests
- Available to biomedical researchers through dbGaP October 1, 2007





NIH GWAS Data Sharing Policy: Guiding Principle

The greatest public benefit will be realized if data from GWAS are made available, under terms and conditions consistent with the informed consent provided by individual participants, in a timely manner to the largest possible number of investigators.

11



Rationale: Sharing Data

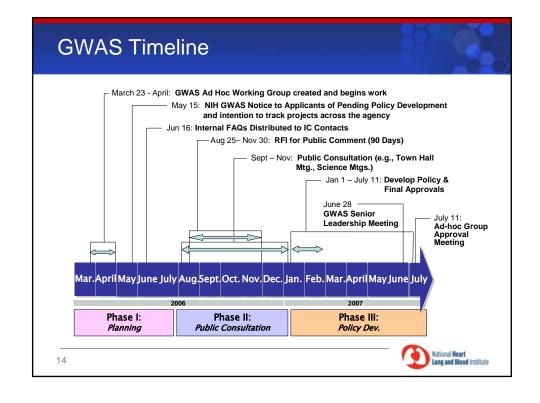
- The opportunities for scientific advances in our understanding of complex, common diseases are now extraordinary.
- The richness of the data generated is far greater than what a single investigator or a group of collaborators can explore
 - Many different scientific questions may be addressed through a single dataset and multiple types of analyses may be needed
 - The opportunity for cross-study analyses increases the capacity to address complex biological questions and can enhance data quality by increasing power
- Information which is not shared represents a lost opportunity to improve the health of the public.
- NIH has been encouraging wide sharing of information for several years.



Why Now?

- The cost of extensive genotyping has fallen rapidly, and continues to fall, making studies feasible which would not have been possible even 4 years ago.
- NIH is receiving many applications for GWAS, representing many millions of dollars of research investment.
- NIH leadership believes that a consistent and robust GWAS policy across the ICs best serves the research community and the public (i.e., research participants)

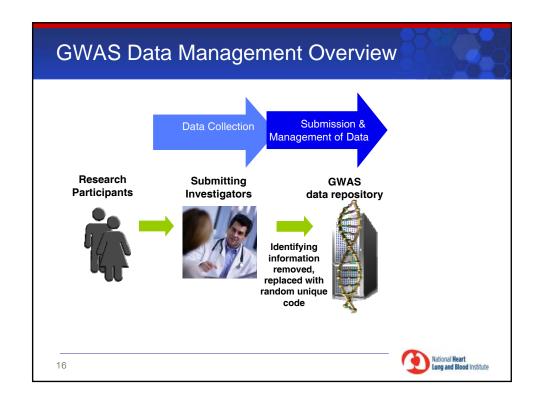




NIH Policy for GWAS

- Data Management
 - Data Submission Procedures
 - Data Access Principles
 - Protection of Research Participants
- Scientific Publication
- Intellectual Property





Data Submission

- Local institution will certify approval of submission to GWAS data repository.
- Certification will include institutional statement that data are provided in accord with all applicable laws and regulations and that an IRB or Privacy Board has reviewed the submission plans.
- Information regarding any limitations on data use is requested at time of application (e.g., limitations imposed by existing informed consent).
- The GWAS Database itself will not be engaging in human subjects research, according to OHRP.
 - Data will be coded by submitting investigators (no HIPAA identifiers)
 - Agreements will be signed stipulating that the identities of research participants will not be disclosed to the NIH GWAS data repository

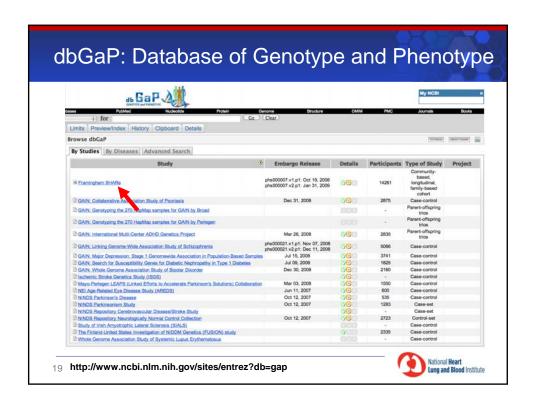
17

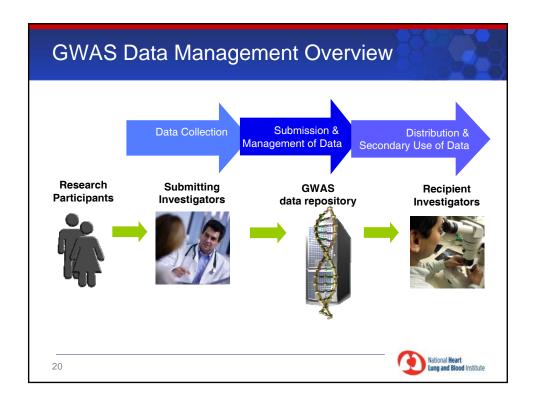


Data Submission - Points to Consider for IRBs

- Intent is to provide investigators and IRBs reviewing datasets for GWAS submission with information on important participant protection considerations
- Topics include:
 - Background on the scientific opportunities presented by GWAS
 - Elements of the GWAS policy
 - Discussion of the ethical issues relevant to the review of submission plans for GWAS datasets
 - Benefits, risks and safeguards that will be used to protect the data
 - Specific points to consider in the evaluation of informed consent documents







Data Access: Data Access Committees (DACs)

- Applications for datasets will be submitted to dbGaP, which then will triage applications to the responsible NIH Institute.
- Each NIH Institute has constituted a DAC for administrative review of applications. DACs are constituted by Institute staff.
- DACs review the application and relay the approval back to dbGaP who releases the dataset.
- DACs consist of Federal staff with expertise in science, bioethics, and privacy/confidentiality issues.

21



Data Access: Data Access Committees (DACs)

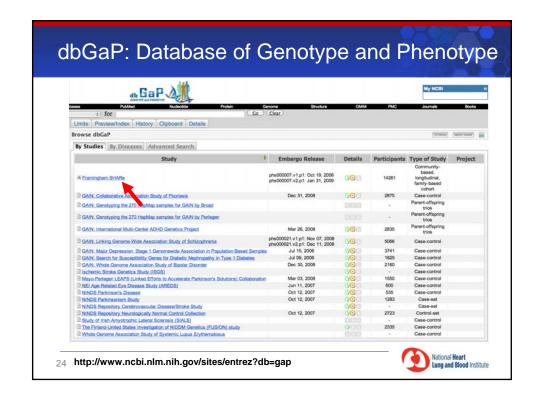
- DACs will determine which investigators will have access to the datasets in the NIH GWAS data repository based on the proposed research use of the data.
- DACs will also play a critical role in protecting the research participants.
- Annual reports from recipient investigators will be required as a condition for continued use of the datasets.
- DACs will review the annual reports for new issues and to confirm that research use matches approved request.



Data Use Certification

- Investigators and their institutions who request GWAS data must provide the following through a Data Use Certification agreement:
 - Description of proposed research projects
 - Agreement to use the data only for the approved research
 - Agreement to follow appropriate data security protections
 - Agreement to follow all applicable laws, regulations, and local institutional policies and procedures for handling GWAS data
 - Agreement not to attempt to identify individual participants within a dataset
 - Agreement not to sell any of the data elements from datasets obtained from the NIH GWAS data repository
 - Agreement not to share with individuals, other than those listed on the request, any of the data elements from datasets obtained from the NIH GWAS data repository
 - Agreement to follow the GWAS policy with regard to publication and intellectual property





dbGaP: Public Site

- Search for studies, protocols, questionnaires
- View phenotype summary data
- View genotype summary data
- View pre-computed or published genetic associations
- Identify studies of interest, learn about their consent conditions, and learn how to apply for data access
- Locate potential collaborators for follow up studies
- No individual data can be viewed

25



dbGaP: Controlled Access Site – Security Measures

- Login/password required to download data on local computer after approval granted by DAC.
- On local computer, each data set is encrypted with a unique password for each Approved User and file.
- PI must use unique password to decrypt file on local machine.
- Consent and terms of use for each data set are included with downloaded files.



Publication

- Period of exclusivity for Primary Investigators
 - Proposed period of 12 months for PIs to publish
 - Exclusivity to apply to any public dissemination of the data or analyses
 - Institutes may elect to shorten this time period
- Acknowledgement of contributing investigators and funding organization

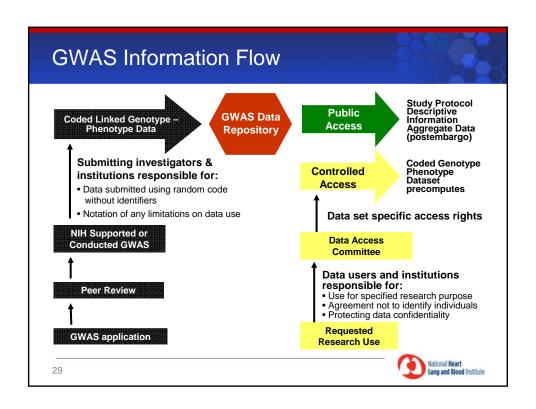
27

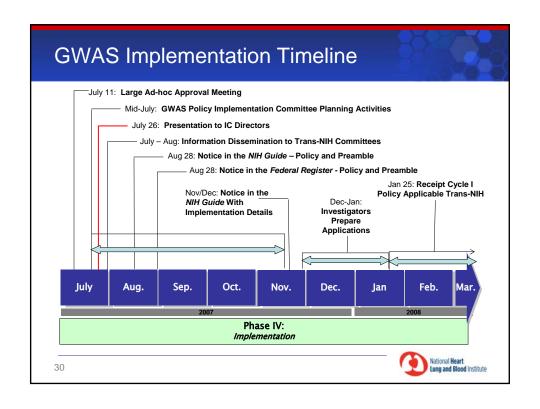


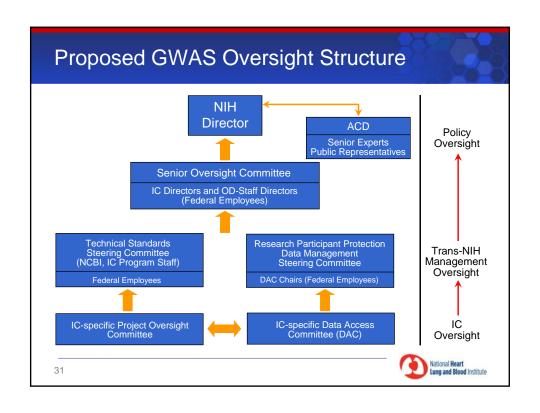
Intellectual Property

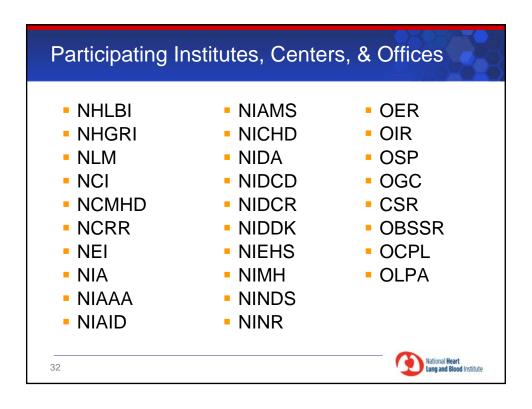
- NIH urges that genotype-phenotype associations remain available to all investigators, unencumbered by IP claims.
- NIH discourages premature claims on precompetitive information.
- NIH encourages broad use of NIH supported genotype-phenotype data consistent with NIH's Best Practices for Licensing with Genomic Inventions.











Acknowledgements

Elizabeth Nabel - Chair Susan Shurin Carl Roth

Christopher O'Donnell Richard Fabsitz Barbara McGarey Valerie Bonham Annette Levey Lauren Higgins Don Schneider Stephen Chanock Deborah Winn Daniela Gerhard Robert Hoover Daniela Seminara Maria Giovanni William Sharrock John Ilekis Jeff Evans Catherine McKeon

Francis Collins Laura Rodriguez Mark Guyer Teri Manolio Lisa Brooks Jean McEwen Elizabeth Thomson Jerome Wilson Katrina Gwinn Anthony Hayward Elaine Collier Hemin Chin James Battey Jerome Wilson Katrina Gwinn-Hardy Anthony Hayward Elaine Collier Zhaoxia Ren Thomas Hart Vivian Ota Wang

Lana Skirboll Marianna Bledsoe Amy Patterson Sarah Carr Norka Ruiz-Bravo Valery Gordon JP Kim Sam Shekar John Burklow Marin Allen David Lipman James Östell Steve Sherry Alan Graeff Michael Gottesman Jerry Menikoff Charlotte Holden Jonathan Pollock Steven Kleeberger Charlene Cho Tom Lehner Melinda Tinkle

