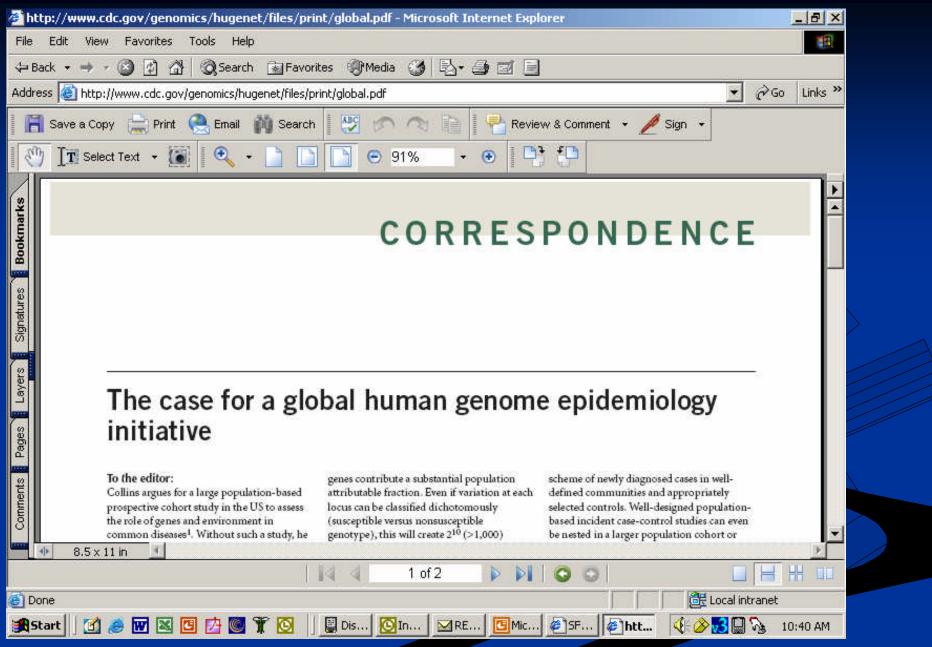
Building the Knowledge Base on Genes and Population Health: Need for Global Collaboration

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Outline: Three Messages

- Global Collaboration in Biobanks/Populationbased Cohort Studies
 - One cohort study in one country is not enough!
- Integration of all Human Genome Epidemiologic Data (cohort, case-control, other)
 - There is more than one way to get there!
- Development of Evidence-based Processes that use Human Genome Epidemiologic Data
 - Direct link from epidemiology to practice!



The Epidemiologic Approach to Genes and Health in Populations: Human Genome Epidemiology (HuGE)

- Gene Discovery: Mostly from family Studies, LD mapping usually from convenient, non-representative groups
- Population-based epidemiologic studies:
 - Gene Variant Characterization: prevalence, genedisease associations in terms of relative, absolute and attributable risks
 - Gene Variant Characterization: gene-gene and geneenvironment interaction



Types of Epidemiologic Studies for Gene Characterization in Populations

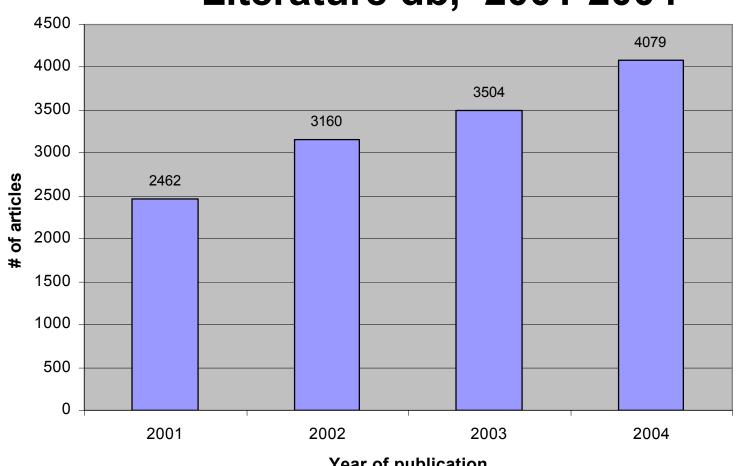
- Cross sectional studies
- Case-control studies
- Cohort studies
- Population biobanks

Myths:

- Stigma around "association studies"
- Cohort inherently superior to case-control studies



No. of articles in Huge Published **Literature db, 2001-2004***



Pub Lit articles

Year of publication

(*As of Jan 14, 2005. Count excludes review articles, meta & pooled analyses.)



Number of Published HuGE Papers* 2001-2004

∠ Year	Prevalence	Associations I	nteractions
z 2001	308	2141	436
z 2002	349	2799	569
z 2003	323	3010	598
z 2004	368	3486	604

•MOST ASSOCIATION STUDIES ARE CASE-CONTROL STUDIES



NHANES III DNA BANK Prevalence of Genes of Public Health Significance

Background

NHANES III DNA Bank

- *National Health and Nutrition Examination Survey (NHANES) is a notionally representative survey
- Detailed interviews, clinical, laboratory and radiologic examinations are conducted
- » Phenotypic data, such as secostatus for many infectious exposures, blood count, chemistres. etc. were collected
- +Ouring second phase NHANES III (1991-1994). white blood cells were hopen and cell lives. were immortalized with EBV
- +NHANES III DNA bank is located at NCEH.CDC. with specimens available from over 7000 participants.
- +in 2002, NCHS announced a call for proposes to use these conceners in the Federal Register.

Challenges to Identifying Genes of Public Health Importance

Selected Pathways of Gene Variants

Numeric Metabolism (e.g., titles and homopystems, lipids, glocose, storbut, stamps to a tempera and inflammatory responses. (e.g., cyloAriela, receptors)
 Activation and inflamosophic perfectly in g., though, temperaports, an accommodal softem perfectly in the commodation and temperaports and the commodation perfectly perfectly selections.

IPAA report perfectly perfectly excellent and (e.g., excellent temperaports) and temperaports. The commodation and temperaports.

Developmental is g., hearing these.

- Goes in information in the tharature

(87 variants of 57 genes)

- Mothocological issues of many available studies
 -Selection basi, power, standards
 Non-restriction of gene-classes assuciation

Collaborative CDC-wide **Proposal Objective**

»Determine the prevalence of genotypes of public health importance.

Laboratory Methods

mong Capametry of Esternal Laboratories to combat the great accombs, true cost, generalizing for HISO DDG SN DI sciences & 87 carteria?

Criteria for Genetic Variants

Public Health Importance

- Known or hypothesized association with diseases of public health importance
- Role in pathways affecting multiple diseases
- » Identified functional variants
- Retailwely common (i.e., ×2.0%).
- » Previously described gene-environment or gene gene interactions
- Resevant phenotypic data available in NHANES dataset
- No current use for clinical risk assessment or intervention.

Public Health Significance of Proposal

- Prevalence of gene variants
 - Basis for estimating population attributable fraction in continuition with measure of gene-disease association
- Enable assessment of potential for screening population
- exceptions for europhility genee. Presidence of constitutions of variants in pollowing and at different local.
- Existing gene-discose association, gene environment and pone-gone interactions

Next Steps

- Frenting against byth NCHS:

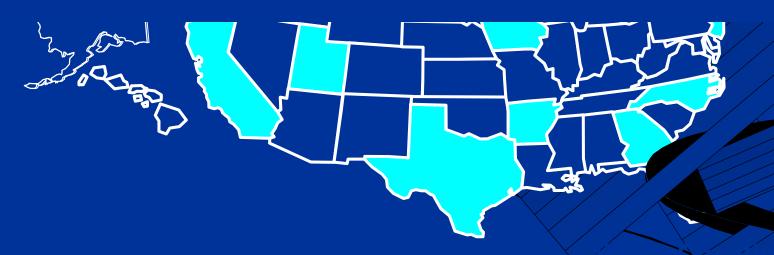
CDC Working Group

ATSDR Clivia Harris. NCBDDD Keren Alse, Cyrrhia Moore, Lorenzo Botte, Quante Yang. NCHSTP. Mary Reschier. NCID. Tom Hodge, Craig Hooper, Jai Lingappa, James McNicoll, Anne Dilley. NCEH. Amenda Brown, Peg Gatlagher, Morta Genn, Omar Henderson, Bruce Lin, Mary Lou Lindegren, Julien Little, Karen Steinberg.
NCCOPHP Hexti Blanck, Wayne Gies, Ingrid Hall, Glusegoina Imperators, Ann Malamber. NIOSH MaryAnn Butler, Annies Weston. PHPPO. Bin Chen. NIP Scott Campbell NCHS. Gerry. McCavillan.





Centers for Birth Defects Research and Prevention



Multi-state Population-Based Case-Control Study Based on State-Based Birth Defects Registries

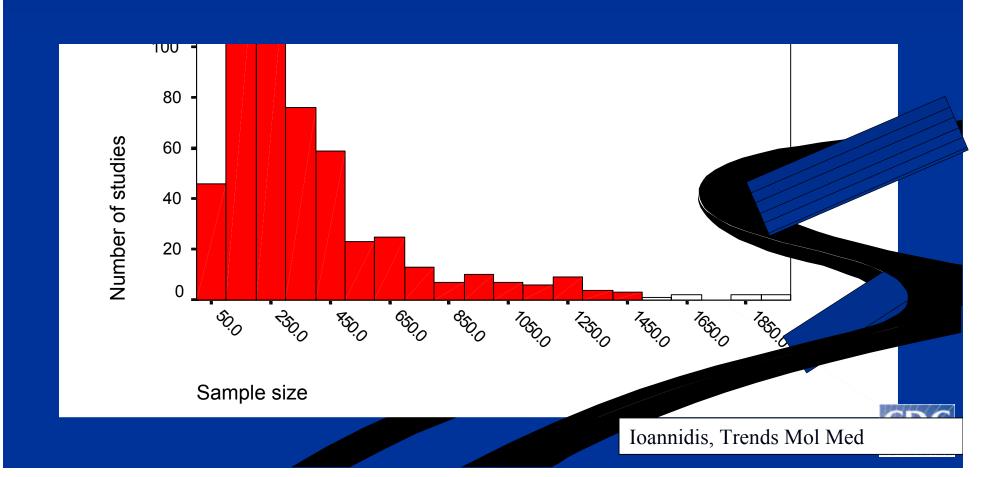


The Need for Integrating Epidemiologic Evidence: on Genes and Population Health

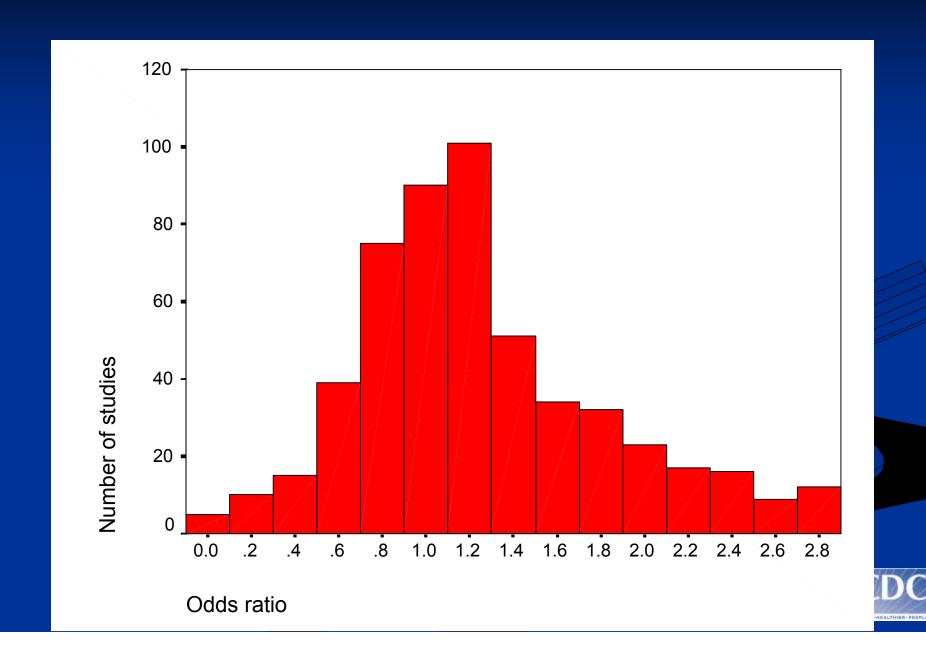
- Unmanageable amounts of data
- Small sample size of individual studies
- Small effect size of gene-disease associations
- Replication of associations
- Publication bias
- Heterogeneity
- Generate and test hypotheses



Small sample size of individual studies



Small effect sizes in individual studies



How to Build the Knowledge Base on Genes and Population Health

- Single large population cohort study
- Systematic synthesis of data from existing and planned cohort studies
- Systematic synthesis of data from all epidemiologic studies (cohorts, case-control, etc)
- Accelerated systematic synthesis of group and individual data using collaborative networks an consortia of all types of studies (cohorts, casecontrol, etc)
- All of the above



Human Genome Epidemiology Network (HuGE Net)

- Global collaboration of individuals and organizations to assess population impact of genomics and how it can be used to improve health and prevent disease
 - Information Exchange
 - Training and Technical Assistance
 - Knowledge Base Development
 - Information
 Dissemination



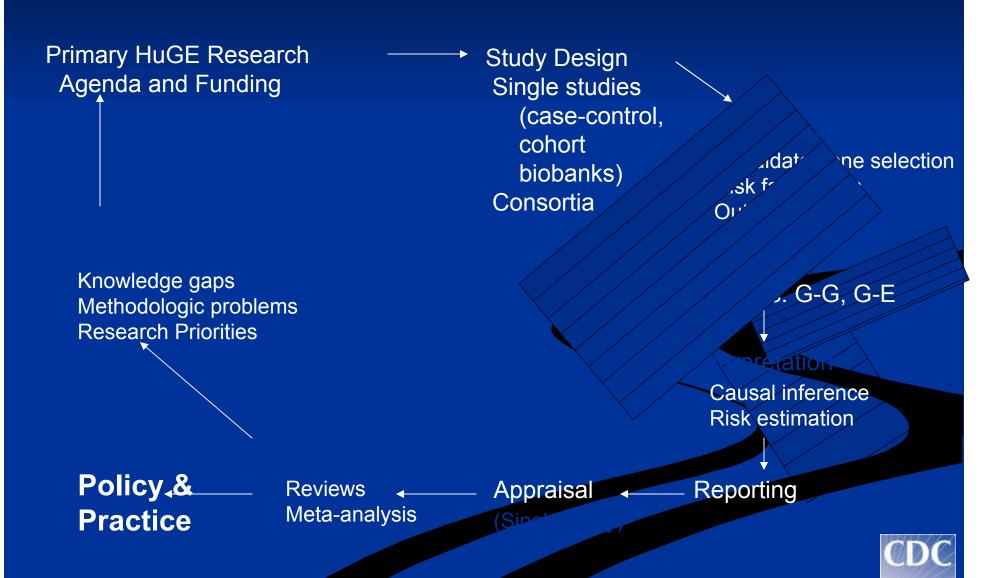


Selected HuGE Net Activities

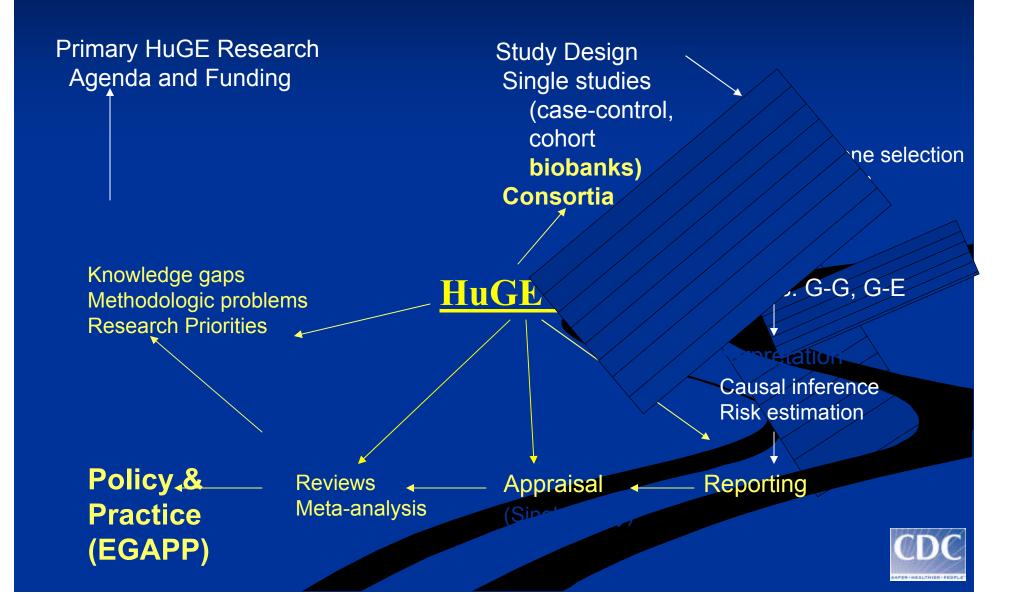
- HuGE Studies Database
- ∠ HuGE Reviews
- Methodology/Training Workshops
- International Biobank/Cohort Study meeting
- "Network of Networks"
- Connecting epidemiologic information to evidence-based evaluation of genomic applications in practice and prevention (EGAPP)

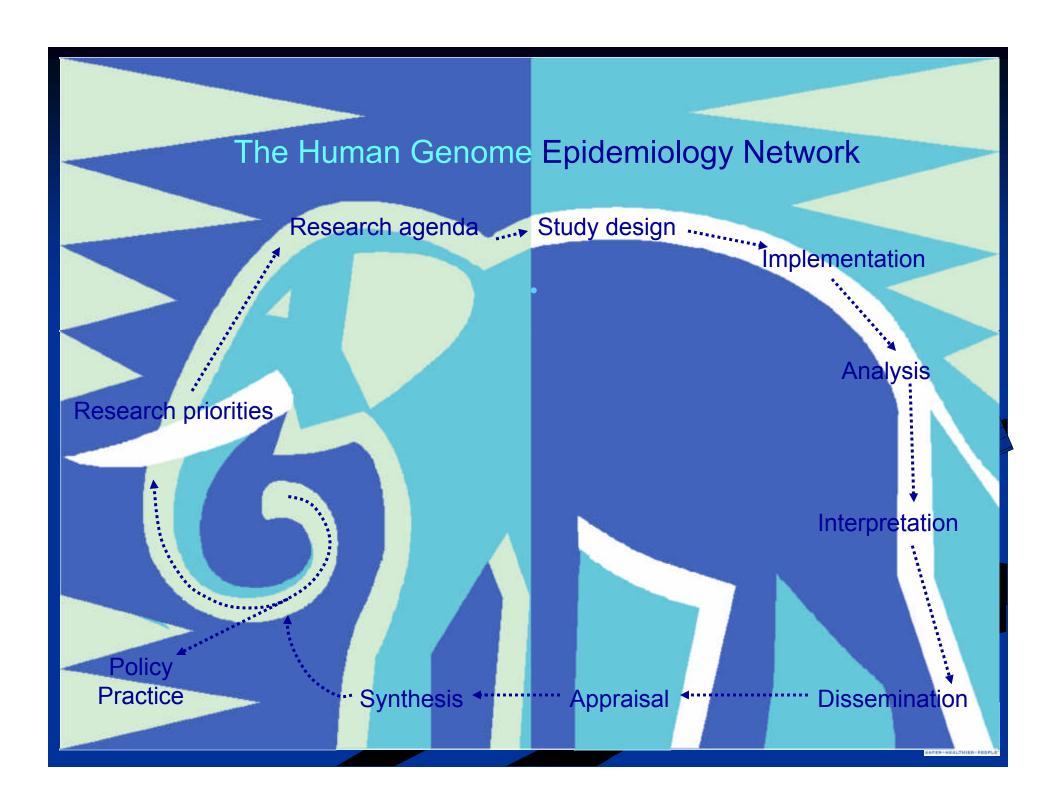


From HuGE Research to Synthesis & Dissemination for Policy and Practice



From HuGE Research to Synthesis & Dissemination for Policy and Practice





International Biobank and Cohort Studies: Developing a Harmonious Approach



Sheridan Buckhead Hotel Atlanta, GA February 7-8, 2005







Biobank Meeting Discussions and Outcomes

- STROBE like statement for publishing studies derived from biobanks
- General report on the design and conduct of biobanks
- Online knowledge base with register of studies and tools (e.g. protocols, informed consent)
- Future periodic meetings with information sharing



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