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# Translation of genomic discoveries to primary care – A role for the PA?

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# Translating Genomics...

- Genomic discoveries relevant to common disease diagnosis and management are coming at an increasing rate.
- Basic discoveries are leading to the development of clinical applications.

## Mind the gap!

- Ergo, improved healthcare is around the corner!



# Translating Genomics...

## Filling the gap

- » Does the application address a clinical need?
- » Does the application meet a clinical need?
- » Is the application acceptable to patients?
- » Is the application acceptable to health care providers?
- » Is the application acceptable to insurers?
- » Is the application acceptable to society?
- » How are patients best educated about the application?
- » How are providers best educated about the application?

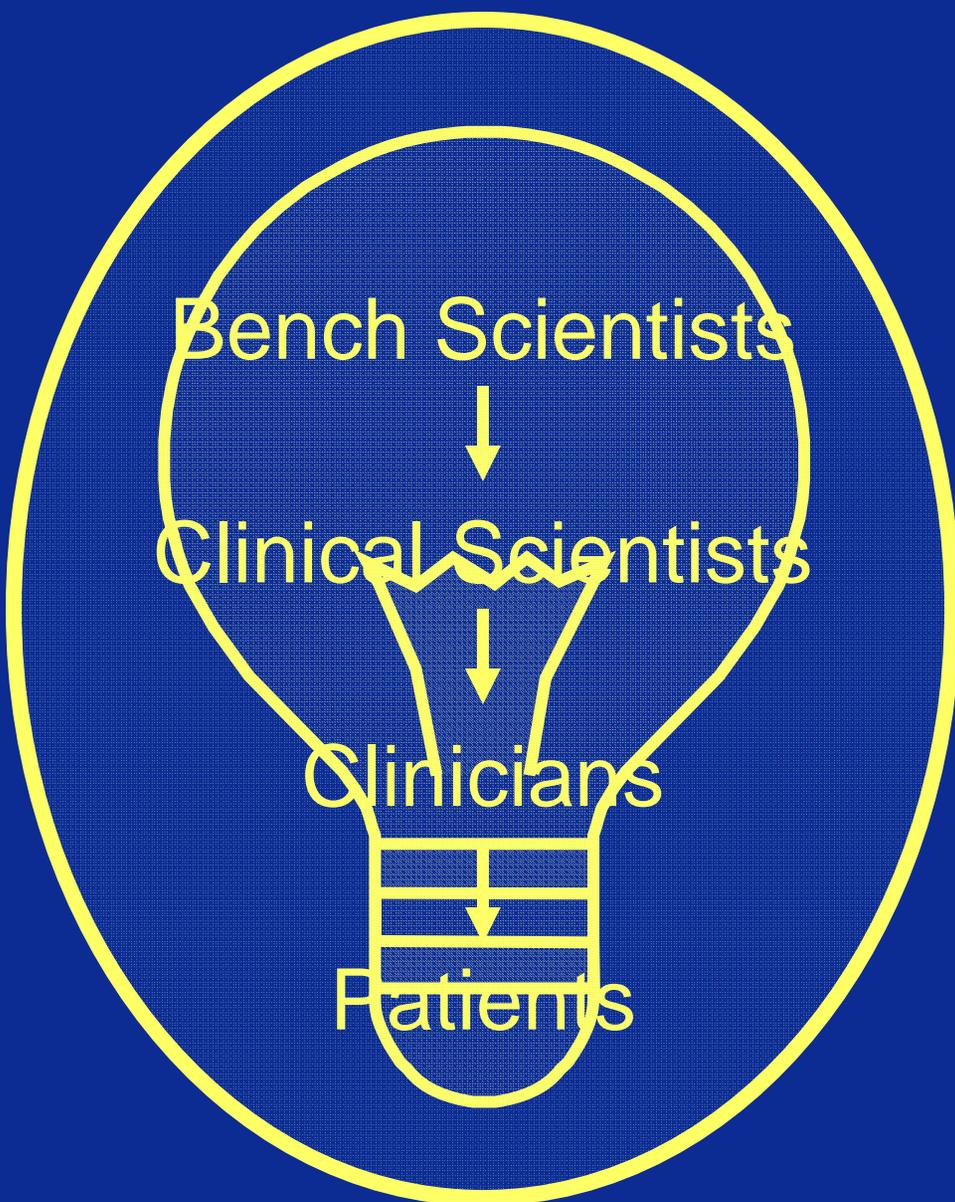


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Who will fill the gap?





# Multiplex ClinSeq

## PUHGV



# Multiplex Genetic Susceptibility Testing:

*A prototype for applied research to inform  
personalized medicine*

**Colleen M. McBride, PhD. & Larry Brody, Ph.D.**

## **Research Partners:**

National Human Genome Research Institute  
Henry Ford Health System  
Group Health Cooperative  
Cancer Research Network (NCI)





# Multiplex Project Aims

## To develop a prototype for multiplex genetic susceptibility testing

- Multiple markers of susceptibility for multiple diseases
- Provide risk feedback to target populations

## To create an infrastructure to facilitate public health research

- Decide upon “standard of care” for consent, feedback & support services
- Identify optimal study population(s) & recruitment approach

# Clinic-based population

## ➤ Cancer Research Network (NCI-funded)

- Full complement of preventive services
- Patient bases geographically distributed with racial-ethnic & SES diversity
- Henry Ford Health System clinical recruitment site
- Group Health Cooperative (HMO Research Network),  
Survey coordination

## ➤ Sample size: 5000+ touched ~ 1000 tested

## ➤ Healthy adults

- Ages 25-40
- Without diseases included on test batter

# Study Design

Baseline screening survey



Mail invitation to website  
to consider genetic testing



Web-based  
decision process re: testing  
w/financial incentives



Consent process  
In-clinic blood draw



Test feedback provided directly to subject  
by mail + telephone follow-up



3 month follow-up telephone survey

# ClinSeq: A translational research project in clinical genomics



Medical & Statistical  
Genetics



NHLBI  
□



NIH Clinical Center



NIH Intramural  
Sequencing Center

# Specific Aims

1. Develop a robust infrastructure for the generation and use of LSMS in a clinical research setting
2. Use LSMS data to develop novel approaches to clinical biomedical research
3. To understand how to interact with subjects re LSMS

# Approach

- Phenotype 1,000 subjects
- Sequence 200-400 candidate genes
- Follow-up studies
- Interpret variants and validate *some*
- Return results

# Clinical evaluation

- Family history (semiautomated)
- Medical history (form-driven)
- Blood pressure
- Coronary calcium score (MDCT)
- Echo/electro-cardiography
- Clinical & research bloods

*Prior to NIH visit:*

- Verbal consent via phone communication
- Family history tool (online)



*Initial visit to NIH:*

- Sample collection for fasting labs (cholesterol, etc)
- General consent
- Family history if unable to complete this information prior to visit
- Medical history intake
- Clinical evaluation
- Second sample collection (non-fasting)



*Visit to Suburban Hospital:*

- Multidetector computed tomography (MDCT) to assess coronary artery calcification



*Initial follow-up (regular mail):*

- Assessment of clinically validated test results (labs, MDCT)



Contact by phone or regular mail to find out if participant is interested in (a) undergoing further phenotyping AND/OR (b) learning genotyping results

AND/OR

Participant may "OPT OUT" of learning results  
AND  
still remain part of study

NOTE 1



*Follow-up visit to the NIH:*  
Genetic education & counseling for results from genome sequencing

*Follow-up visit to the NIH:*  
Further phenotyping

Health Professionals' Genetics Education Needs Exploration (HP GENE) Survey



National Human Genome Research Institute

National Institutes of Health

# Health Professionals' Understanding of Human Genetic Variation Study

Vence Bonham, JD  
Associate Investigator  
Social and Behavioral Research Branch  
Principal Investigator

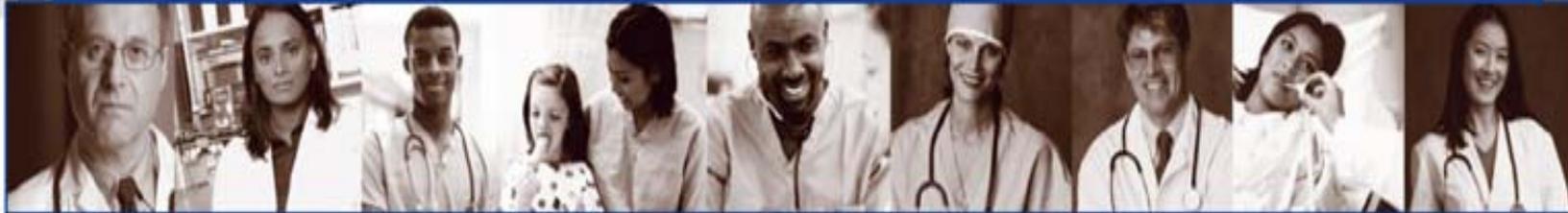


# Project Aim

To investigate health professionals' **knowledge** of human genetic variation, **beliefs** about biological and genetic differences based upon their patients' race and ethnicity and its **use** in clinical practice.



## Health Professionals' Genetics Education Needs Exploration (HP GENE) Survey



National Human Genome Research Institute

National Institutes of Health



**7. Random mutations cause all of the genetic variation in the human genome.**

- |                       |                       |  |                       |
|-----------------------|-----------------------|--|-----------------------|
| true                  | false                 | scientific<br>evidence<br>inconclusive | don't<br>know         |
| <input type="radio"/> | <input type="radio"/> | <input type="radio"/>                  | <input type="radio"/> |

**8. The variation in the human genome includes both disease causing gene variants and variants that have no effect on health and disease.**

- |                       |                       |  |                       |
|-----------------------|-----------------------|--|-----------------------|
| true                  | false                 | scientific<br>evidence<br>inconclusive | don't<br>know         |
| <input type="radio"/> | <input type="radio"/> | <input type="radio"/>                  | <input type="radio"/> |

## STUDY DESIGN

### **Phase I Qualitative Study**

Dimensional analysis and qualitative content analysis were used to explore physicians' perceptions of and experiences with racial factors in clinical decision-making, determining the racial background of a patient, and perceptions of the race-related causes of health differences.

### **Phase II Scale Development**

Focus groups were used to assist in question development. The process of scale development occurred in an iterative fashion. Thirty-two cognitive interviews with physicians were used to refine the instrument and scale. Two panels of experts, geneticists with expertise in human genetic variation and social scientists with expertise in survey methodology provided input.

### **Phase III National Physician Survey**

A pilot survey of 400 physicians will be conducted fall 2007 to examine psychometrics of the scale. The scale will be revised based upon the findings. In 2008 a National Survey of 3000 Primary Care Physicians will be conducted using the final HGVB scale.

### **Phase IV National Physician Assistants Survey ????**





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**ClinSeq**
- Vence Bonham, J.D., ECIB, NHGRI  
**PUHGV**



# Possible discussion topics:

- To what extent will these sorts of research questions interest the PA community?
- What unique perspectives could the PA community bring to this type of research?
- To what extent do PA training centers participate in research? Independent? Part of a larger academic center?
- Do PA's have a research society? NAPCRG? How to engage PA's with interest?



# Possible discussion topics:

- What other factors need to be considered to facilitate the translation of genomic discoveries to primary care?