

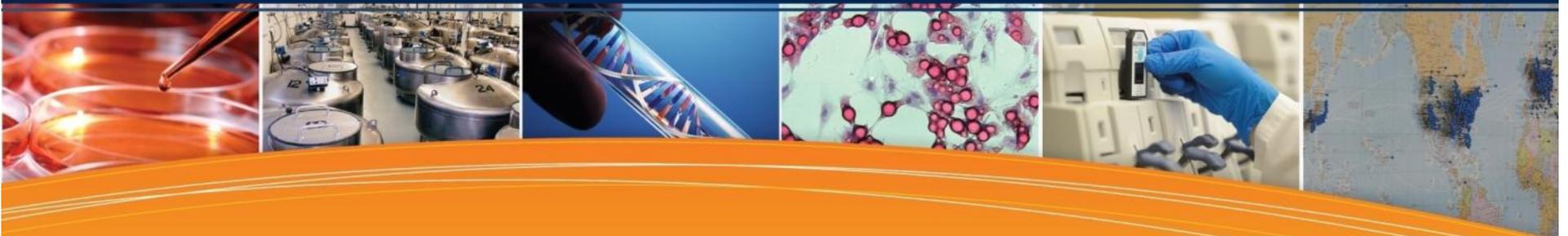


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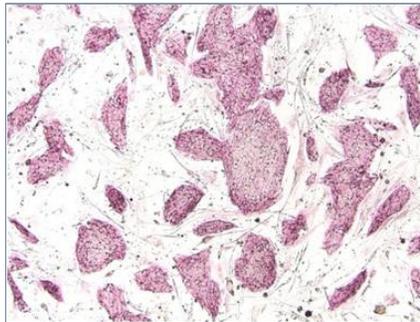
FOR MEDICAL RESEARCH

Utility of Genome Information in Clinical Care: *Coriell Personalized Medicine Collaborative*[®]

Michael Christman, Ph.D.
President & CEO, Coriell Institute
July 19, 2011



Coriell Institute for Medical Research



CPMC Research Study Launched in 2007



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Goals of the CPMC Research Study

- *Study the use of genome-informed medicine in a real-world clinical setting*
- *Determine the best mechanism to provide information to providers and participants/patients*
- *Find correlations in observational data*

www.cpmc.coriell.org



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Dr. Francis Collins

Interview with *Science Magazine*

June 6, 2008



*“We desperately need, in this country, a **large-scale, prospective, population-based cohort study**. And we need to enroll at a minimum half a million people. We would need to have their environmental exposures carefully monitored and recorded, their DNA information recorded, their electronic medical records included, and have them consented for all sorts of other follow-ups.”*



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How the CPMC Study works



1

My Clinical Data

Why do I need to give this information?
The answers you provide in the questionnaire below will be used to develop personalized risk reports. The answers you provide in these questionnaires for the study will not be able to produce your results. To make sure we are providing accurate risk results we will ask you to update your information on a yearly basis.

You do not need to complete all of the questionnaires at one time. You can save your answers at any point and return at a later time.

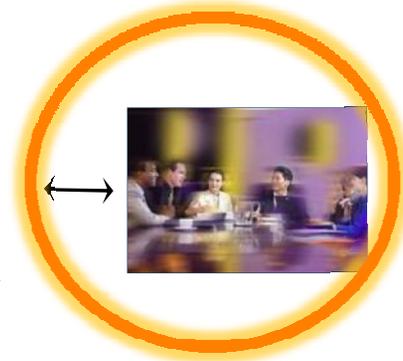
Questionnaire	Estimated Time	% Complete
Demographics	5 minutes	100%
Family History	40 minutes	75%
Lifestyle	5 minutes	100%
Medical History	20 minutes	100%
Medications	5 minutes	100%
Interactions with Healthcare	5 minutes	100%
My Questionnaires		
Current Knowledge Number	30 minutes	100%
My Genetic Results		

2



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3



5



4



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Hospital Partners in the CPMC

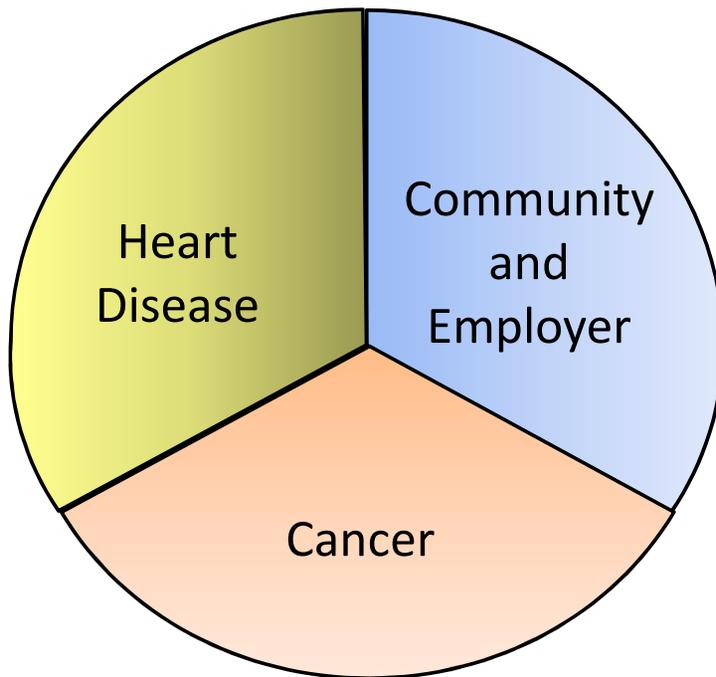


CPMC Recruitment



Eligibility:

- *Be at least 18 years old*
- *Hear informed consent presentation*
- *Have an email address and access to the internet*



Recruitment Mechanisms:

- *Community – based*
- *Cancer Clinic – based*
- *Heart Clinic– based*



What information do we collect?

All Cohorts:

Demographic Information
Medical History
Medications
Family History
Lifestyle Information
Genetic Knowledge assessment



Cancer Cohorts:

Cancer Registry Data
Cancer-related health records
Prescribing Records



Heart Disease Cohorts:

Electronic Health Records
Prescribing Records



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Detailed:

Family, Medical History, Medication, Lifestyle, and Demographic Information

My Clinical Data

Why do I need to give this information?

The answers you provide to the questionnaires below will be used to develop personalized risk reports. The sooner you complete these questionnaires the sooner we will be able to process your results. To make sure we are providing accurate risk results we will ask you to update your information on a yearly basis.

You do not need to complete all of the questions at one time. You can save your answers at any point and return at a later time.

		Estimated Time Req.	% Complete
Demographics		5 minutes	 100%
Family History		45 minutes	 8%
Lifestyle		5 minutes	 100%
Medical History		20 minutes	 0%
Medications		5 minutes	 100%
Medications with Reactions		5 minutes	 100%

My Questionnaires

		Estimated Time Req.	% Complete
Genetic Knowledge Review		30 minutes	 100%

My Genetic Results



The CPMC uses two “GeneChips”



2 million sites of variation

*2,000 sites of known
relevance to drug action*



Who decides what genetic information is reported?

- *Informed Cohort Oversight Board (ICOB), an external advisory board. Composed of scientists, medical professionals, ethicist, community members.*



- *Vote on whether conditions are potentially actionable.*
- *Meet at least twice a year to approve new conditions.*
- *New results then reported to ALL participants.*

Supported by RNR Foundation



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Informed Cohort Oversight Board (ICOB)

Robert C. Green, MD, MPH

Boston University, MA

Steven A.R. Murphy, MD

The Personalized Medicine Group, CT

Erin O'Shea, PhD

Harvard University, MA

David Pellman, MD

Harvard Medical School, MA

Charles Rotimi, PhD

National Human Genome Research Institute, MD

Reverend Floyd White

Woodland Community Development Corp, NJ

Jennifer Hoheisel, MS

Camden County College, NJ

Ellis J. Neufeld, MD, PhD

Children's Hospital Boston, MA

Marc Lenburg, PhD

Boston University School of Medicine, MA

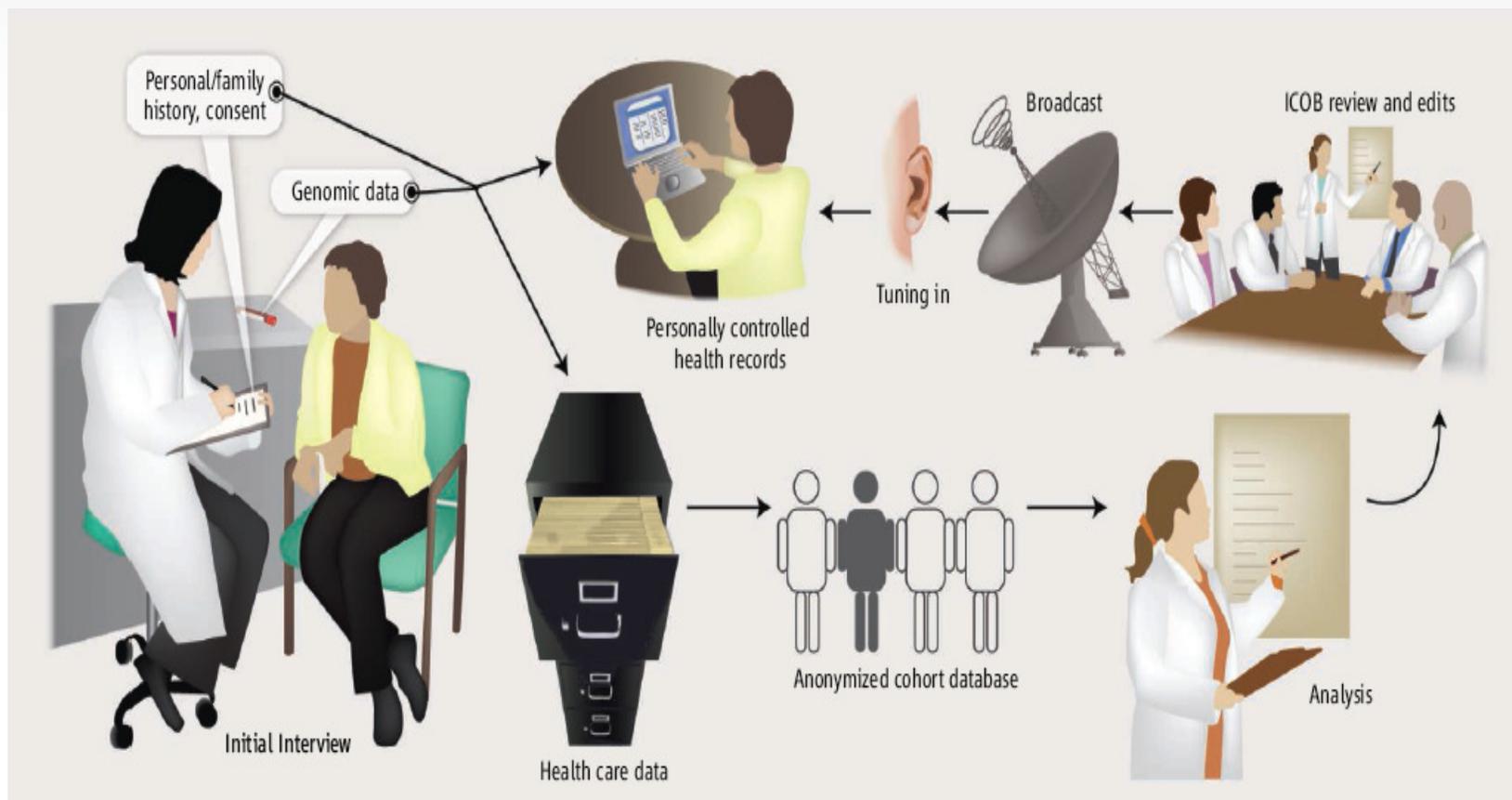
Kenneth Offit, MD, MPH

Cornell University, NY



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Informed Cohort Concept

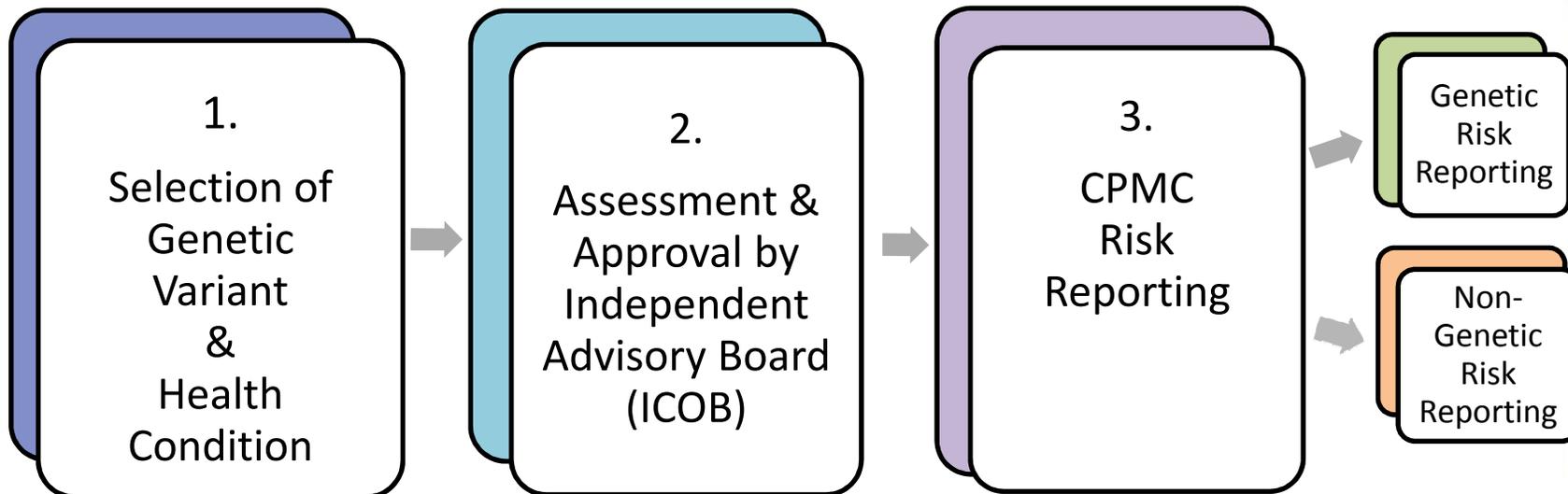


Kohane et al., Science 2007



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CPMC Process Overview



Potentially Actionable Conditions Currently Approved to be Reported by the CPMC Study®

Drug Metabolism

CYP2D6

VKORC1

CYP2C9

CYP2C19

UGT1A1

CYP4F2

Complex Disease

Age-related macular degeneration

Breast cancer

Bladder cancer

Chronic obstructive pulmonary disease

Colon cancer

Coronary artery disease

Inflammatory bowel disease

Hemochromatosis

Lupus

Melanoma

Obesity

Prostate cancer

Rheumatoid arthritis

Testicular cancer

Type 1 diabetes, and Type 2 diabetes



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My New Genetic Results

To view your PERSONAL result, click “View My Result.”

Type 2 Diabetes rs7754840	View My Result	Do Not View My Result	Talk To Genetic Counselor First
Prostate Cancer rs16901979	View My Result	Do Not View My Result	Talk To Genetic Counselor First
Age-Related Macular Degeneration rs10490924	View My Result	Do Not View My Result	Talk To Genetic Counselor First
Melanoma rs910873	View My Result	Do Not View My Result	Talk To Genetic Counselor First
Iron Overload/Hemochromatosis rs1800562	View My Result	Do Not View My Result	Talk To Genetic Counselor First

Demonstrating Variants to CPMC Participants



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[About](#) | [Results](#) | [Limitations](#) | [What do I do now?](#) | [Methods](#)

[Genetic Variant](#) | [Risk Summary](#) | [Genetic Variant Risk](#) | [Family History Risk](#) | [Other Risk Factors](#)

Coronary Artery Disease – Variant #1 (rs1333049)

We all have 2 copies of every gene, one from each of our parents.
Each copy may have small changes called genetic variants.
Some genetic variants are associated with an increased risk of disease.

Your Result: 1 copy of the risk variant and 1 copy of the non-risk variant were detected (GC)

Details

Name of Gene:

CDKN2A/CDKN2B
on chromosome 9p21.3

Two Possible forms of Variant 1 (rs1333049):

Risk variant = C
Non-risk variant = G

This variant accounts for only a **small fraction** of the total genetic risk of Coronary artery disease.

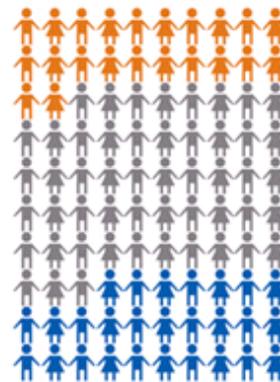
Population Data

CC - 22 in 100 people have 2 copies of the risk variant

GC - 51 in 100 people have 1 copy of the risk variant and 1 copy of the non-risk variant

GG - 27 in 100 people have 2 copies of the non-risk variant

These results are based on studies in Caucasian populations.



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Genetic and Non-Genetic Risk Assessments: *Provided via CPMC Online Web Portal*

Coronary Artery Disease – Variant #1 (rs1333049)

Risk Summary

This graph provides a summary of the Relative Risk for Genetic variant, Family History, and Other Risk Factors.

Place mouse over colored disk for risk information.

Your Risk due to:



Family History



Other Risk Factors

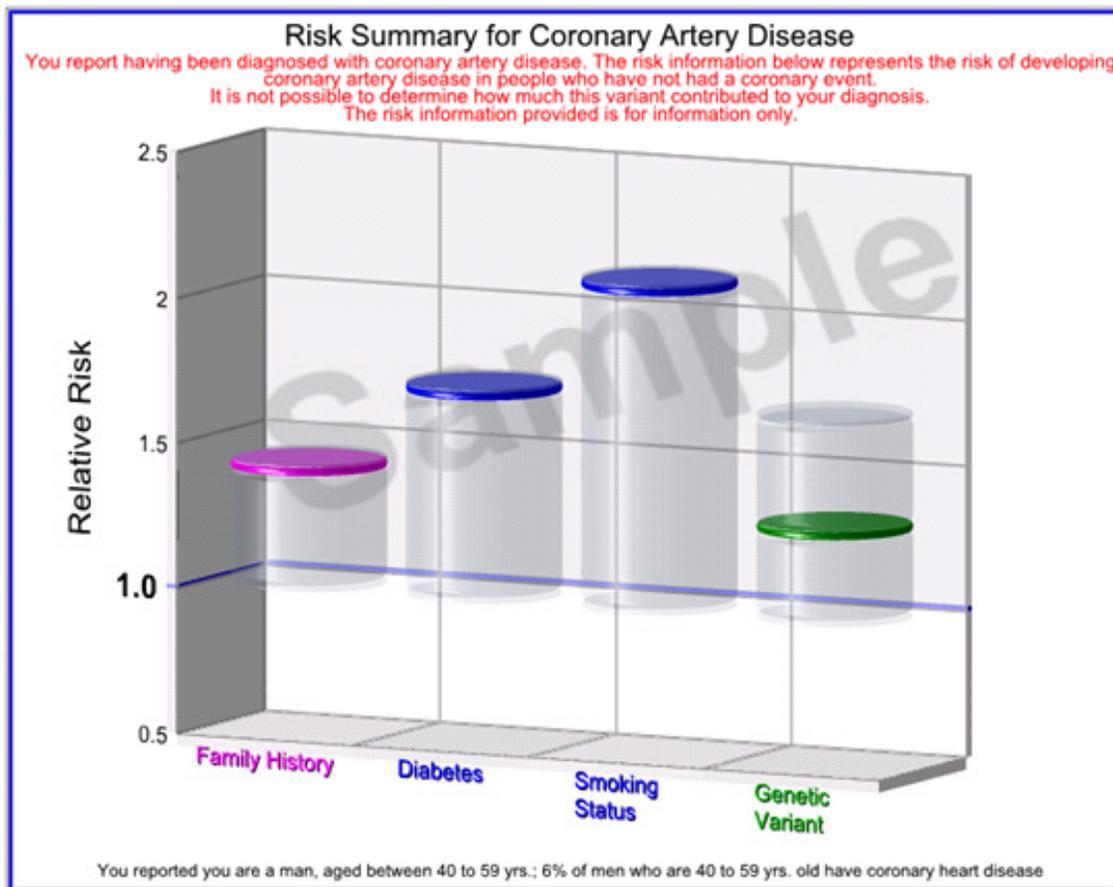
Diabetes
Smoking Status



Genetic Variant



Click on each cylinder for bigger view and more information



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Coronary Artery Disease Surveys: *Provided via Portal at 3 and 12 months*



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how it works

[Informed Consent](#)

[Saliva Collection](#)

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[Genetic Results](#)

[Sharing My Results](#)

[Outcomes Research](#)

Outcomes Research



Your Ongoing Participation is Important to Us

The CPMC research study is a longitudinal study that will continue for a minimum of 5 years. This means that we will stay in contact with you as the study goes on to gather more data. Each year, you will be asked to update your medical history, family history and lifestyle information.

In addition, a few months after you view your genetic variant results, we will ask you to provide feedback through the web portal on whether and how you used the information.

Why are we asking so many questions? The goal of the CPMC™ is to understand how you use your personalized genetic information and if you use this knowledge to improve your health. By telling us whether and how you used your genetic results, you will help us achieve this goal.



So, from the CPMC™ team, thanks for your participation in this important project!



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Coronary Artery Disease: Outcome Survey

Respondent Characteristics	N = 472	%
Mean Age 51 years		
Female	315	68.9
Caucasian	433	94.8
Bachelors Degree or higher	328	71.7
Occupation		
Healthcare providers (all)	122	26.7
Life, Physical, Social Scientists	24	5.3
All other Occupations	311	68.1



Coronary Artery Disease: Outcome Survey

...cont'd:

Distribution of Participants by Number of CAD Risk Factors	N = 430	%
No Risk Factors	55	12.8
One Risk Factors	197	45.8
Two Risk Factors	161	37.4
Three - Four Risk Factors	17	4.0
Participants reporting somewhat/very high perceived risk		
No Risk Factors	1 / 55	1.8
One Risk Factor	63 / 197	32.0
Two Risk Factors	75 / 161	46.6
Three to Four Risk factors	14 / 17	82.4



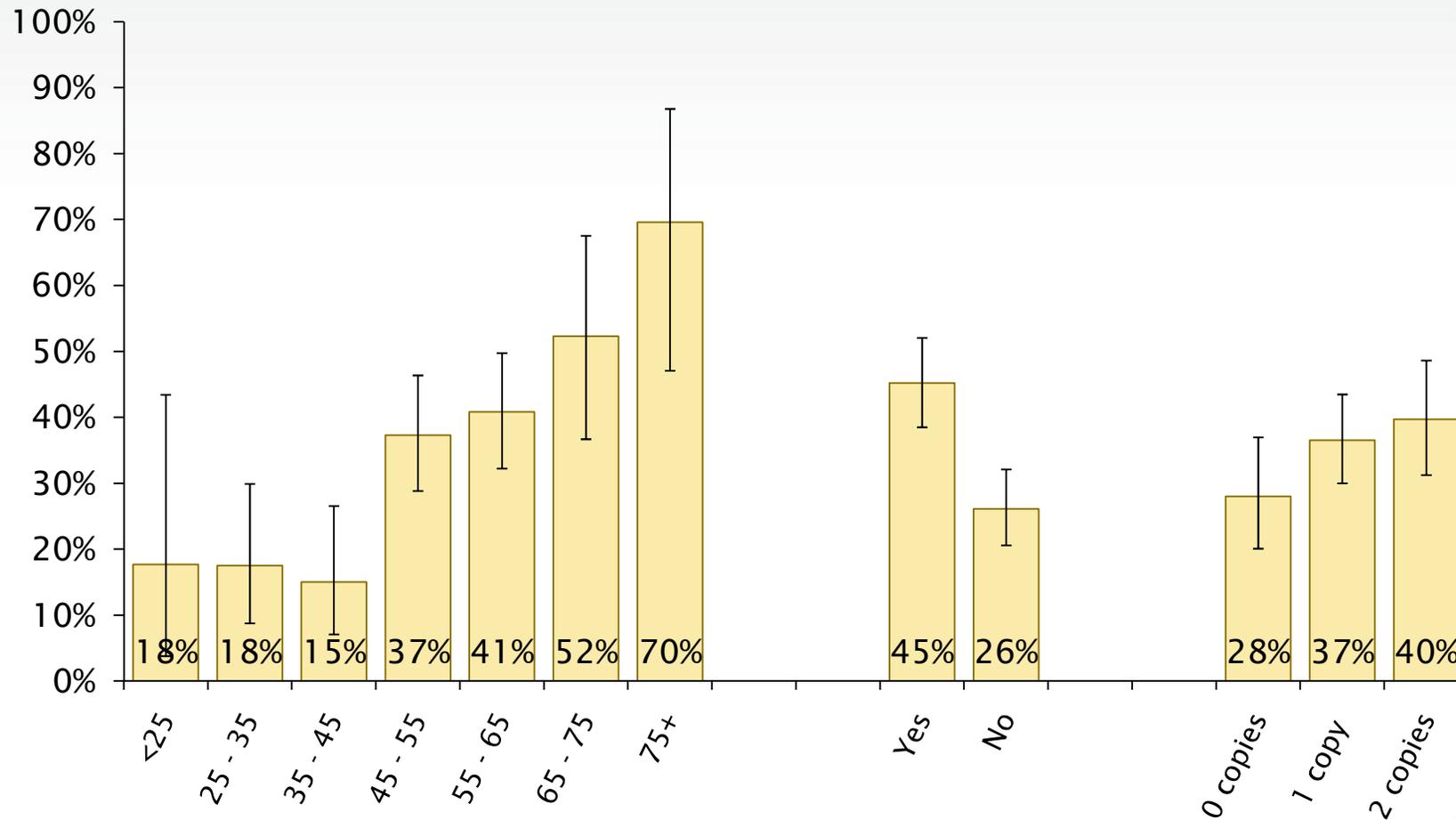
Outcomes by Risk Group

Tests/Procedures received since receiving CPMC CAD results:

Electrocardiogram	95	20.8%
Echocardiogram	42	9.2%
Stress Test	31	6.8%
Nuclear Stress Test	17	3.7%
Electron Beam CT	5	1.1%
MRA	4	0.9%
Balloon angioplasty and stent placement	2	0.4%
Other	34	7.4%
Don't Know	2	0.4%
No tests or procedures	296	64.8%



Tests or Procedures by Risk Group



Tests received associated with age and positive family history of CAD. No strong association between genotype and tests received.



Members:

Pharmacogenomics Advisory Group (PAG)

Marialice Bennett, BS, RPh
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Art Caplan, Ph.D.
University of Pennsylvania, PA

**Michael D. Ezekowitz, MBChB, DPhil,
FRCP, FACC**
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Ohio State University College of Medicine, OH

Issam Zineh, PharmD, MPH
US Food and Drug Administration, MD



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Strength of Evidence Code

Table 1. PGx gene haplotype category assignments based on strength of evidence for drug interaction phenotype

Evidence type	Evidence code	Category type
<i>In vivo</i> clinical outcome for reference drug	1	include
<i>In vivo</i> PK/PD for reference drug	2	include
<i>In vitro</i> enzyme activity for reference drug	3	include
<i>In vitro</i> enzyme activity with probe substrate <u>plus</u> mutation type (n, sb, se, ca)	4n or 4scd or 4se or 4ae	include
<i>In vivo</i> clinical outcome with another drug <u>plus</u> mutation type (n, sb, se, ca)	5n or 5scd or 5se or 5ae	include
<i>In vivo</i> PK/PD for another drug <u>plus</u> mutation type (n, sb, se, ca)	6n or 6scd or 6se or 6ae	include
<i>In vitro</i> enzyme activity with another drug <u>plus</u> mutation type (n, sb, se, ca)	7n or 7scd or 7se or 7ae	include
<i>In vitro</i> enzyme activity with probe substrate <u>only</u>	8	exclude
<i>In vivo</i> clinical outcome with another drug <u>only</u>	9	exclude
<i>In vivo</i> PK/PD for another drug <u>only</u>	10	exclude
<i>In vitro</i> enzyme functional (protein stability or enzyme activity with another drug) <u>only</u>	11	exclude
<i>In vitro</i> or <i>in vivo</i> data does not support functional role	12	exclude
No <i>in vitro</i> or <i>in vivo</i> data	13	exclude
Genotype frequency data suggestive of "private mutation". A private mutation is defined as a genetic variant found in a single individual or single family without being observed in reference populations.	14	exclude

PK=pharmacokinetic;
 PD=pharmacodynamic;
 n=null mutation;
 scd=mutation located in known important substrate-binding or catalytic domain;
 se=mutation leading to splicing error;
 ae=mutation leading to altered gene expression.

- Review *in vivo* PK/PD evidence or *in vitro* functional analysis supporting effect on protein function (e.g. enzymatic activity, plasma concentrations, etc.), if available.
- Review clinical outcome data supporting drug metabolizing phenotype, including adverse events or reduced efficacy.



American Pharmacists Association



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Minority Participation and Outreach

United States Senator Robert Menendez



Senator Menendez delivers remarks at the Coriell Institute for Medical Research to encourage New Jerseyans to participate in this groundbreaking genome research project which aims to benefit individuals suffering from diseases.



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Potential CPMC Study Outcomes

- *Participant Behavior*
- *Physician Behavior*
- *Adverse Drug Reactions*
- *Pharmacogenomic Efficacy*
- *Clinical Outcomes*



CPMC Partnerships and Collaborations



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Baseline Genetic Knowledge: *Survey Results*

- *15 Questions*
- *2,189 participants completed survey*
- *Mean of 76% correct answers*
- *ANOVA model adjusted for multiple participant*
- *Characteristics shows:*
 - *Age inversely associated with correct responses*
($p < 0.001$)
 - *No association between income and correct responses*
($p = 0.74$)



Ancillary Studies

*CPMC Participant Behavior Upon Receiving Genome Info
(Barbara Bernhardt and Reed Peyritz at Penn)*

- **Assess motivations to participate** in the CPMC and perceptions of the utility of personalized medicine

Recruit potential “early adopters” to complete an anonymous survey from among those who sign up to attend a CPMC enrollment event

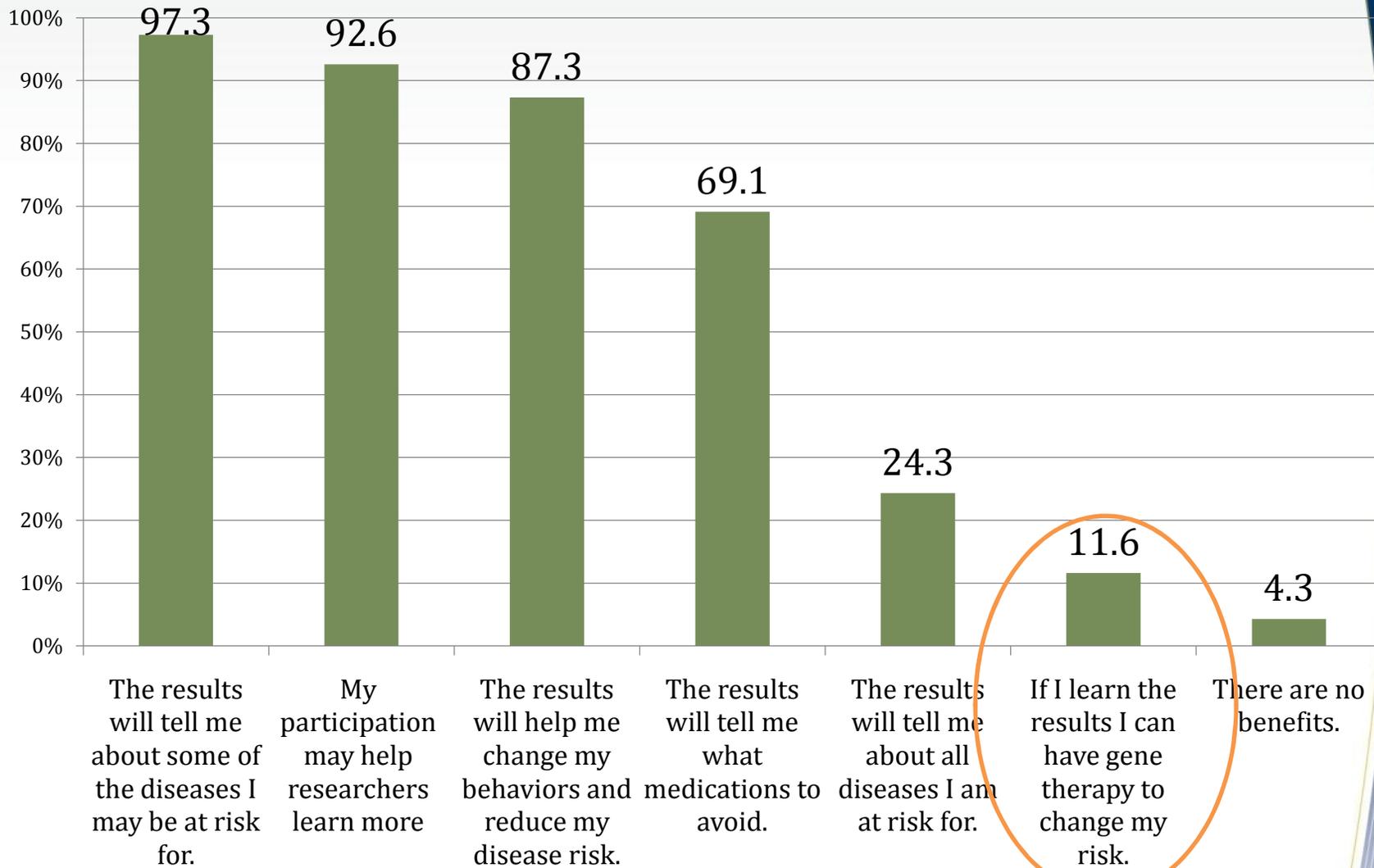
- **Explore participant understanding** of personalized genomic disease risk results, intended and actual use of information and educational needs of individuals receiving results

Interview with CPMC participants (n=60)

- **Develop possibly recommendations** for the ethical offering of personalized genomic disease risk assessment



Perceptions of Benefits of CPMC

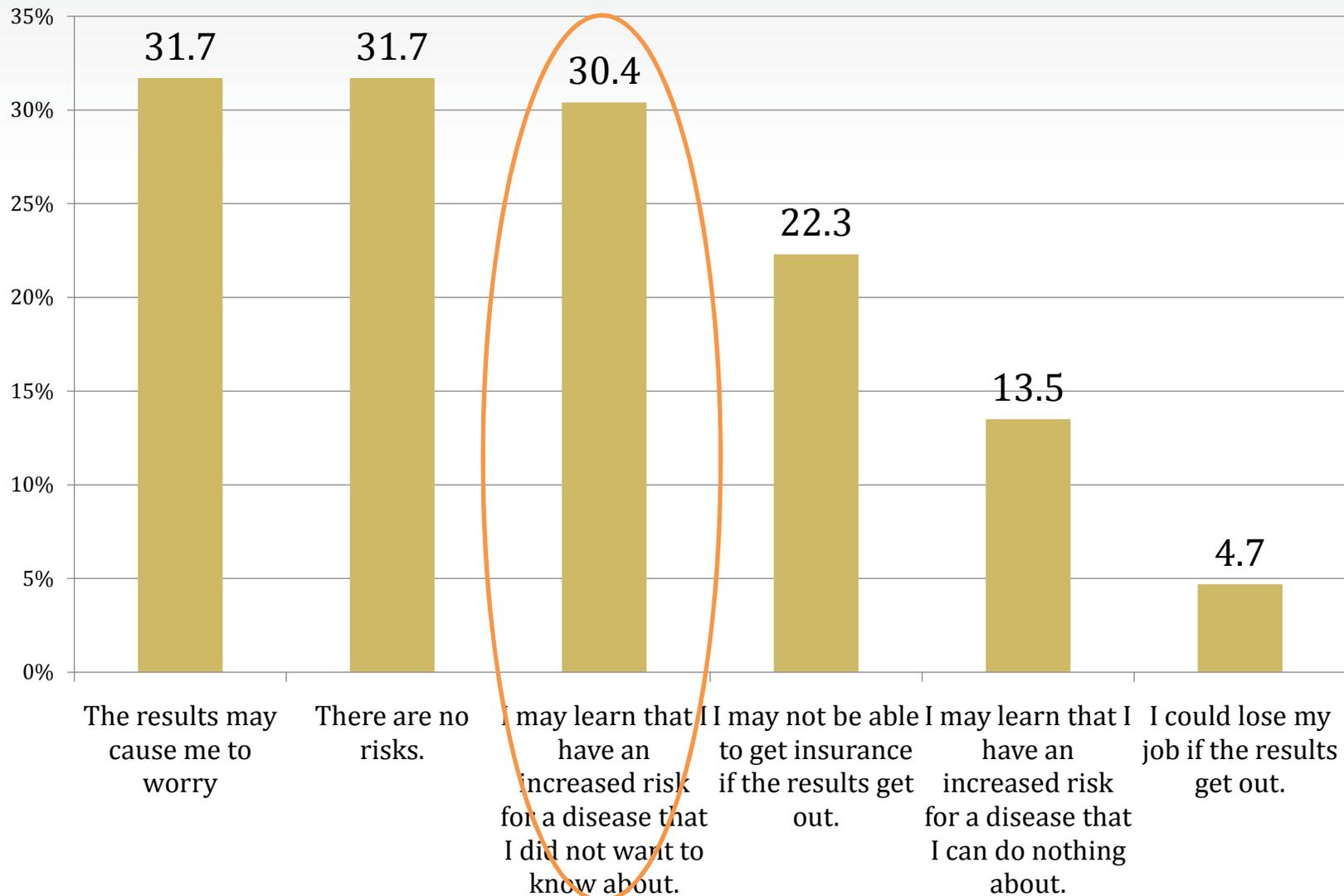


% agreeing or strongly agreeing



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Perceptions of Risks of CPMC



% agreeing or strongly agreeing



CPMC at OSUMC



- Patient Participants (n=1800)
 - *Congestive heart failure patients*
 - *Hypertensive patients*
- Physician Participants (n=30)
 - Cardiologists
 - Primary care physicians



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CPMC at OSUMC



- *Study Design (patients)*

Patients will be identified through enrolled physicians

Patients who consent will complete MFLQ and baseline assessment (risk perception, numeracy assessment, etc)

Patients will be randomized to in person genetic counseling or no counseling (participants in no GC arm will have access to CPMC genetic counselors for urgent questions)

Follow up assessment of risk perception, understanding of results, satisfaction, information seeking, etc

Analysis of impact of genetic counseling in genomic testing



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CPMC at OSUMC



- *Exploratory Aims (Pharmacogenomics)*

To determine if genetic data are effective at differentiating congestive heart failure (CHF) responders from non-responders

To compare disease progression/negative outcomes between CHF responders and non-responders

To determine if genetic data are effective at predicting which newly diagnosed hypertensive patients will require modifications and/or additions to their initial treatment



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Observational Data Can Be Useful



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Exploring Best Practices:

Guiding the ethical, legal and responsible implementation of personalized medicine

- *Study participants control information*
- *Genetic counseling offered*
- *Report quantitative non-genetic risk*
- *Web portal provides two way communication with participants*
- *ICOB provides dynamic reassessment of genomic data*
- *Report on only “potentially actionable” conditions*
- *Seek expert advice on actionability (ICOB, PAG)*



Big Picture: *Ethical, Legal and Social Issues*

- *Ensuring genetic privacy*
- *Reducing anxiety associated with genetic prognosis*
- *Updating medical record technology*
- *Educating the community, doctors, nurses, pharmacists, and genetic counselors*
- *Payors likely to drive clinical adoption*



MIT Technology Review 2010



Top 10 "Research To Watch"

April 2010



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