

## Scientific Evidence for Personalized Nutrition: Ethical Implications of Methodological Limitations

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# A Critical Appraisal of the Scientific Basis of Commercial Genomic Profiles Used To Assess Health Risks and Personalize Diet and Lifestyle Interventions

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Erasmus University Medical Center Rotterdam, Rotterdam, the Netherlands Centers for Disease Control and Prevention, Atlanta GA, USA

EGAPP, February 2008, Atlanta GA



## Premature applications of genomic profiling





#### CELLF COMPREHENSIVE

A combined analysis of nineteen genes that may play an important role in how your body manages bone health, heart health, antioxidant and detoxification function, insulin sensitivity and inflammation. Perfect for those who want a complete snapshot of their overall health profile and recommendations for achieving optimal health without a specific disease focus.

Price: \$252.00

### **Profiles**

### Disease-specific

- Heart health or cardiogenomic profile
- Bone health or osteogenomic profile
- Inflammation health
- Estrogenomic
- Immunogenomic

### General

- Enhanced Basic Screening Panel
- Genoscore
- Nutritional Genetic Profile
- Personal DNA Analyses Starter Kit
- ٠...



### **ARTICLE**

A Critical Appraisal of the Scientific Basis of Commercial Genomic Profiles Used to Assess Health Risks and Personalize Health Interventions

A. Cecile J.W. Janssens,<sup>1,\*</sup> Marta Gwinn,<sup>2</sup> Linda A. Bradley,<sup>2</sup> Ben A. Oostra,<sup>3</sup> Cornelia M. van Duijn,<sup>4</sup> and Muin J. Khoury<sup>2</sup>

The American Journal of Human Genetics 82, 593–599, March 2008 593

METHODS: Reviewing meta-analyses of genetic associations for the genetic variants included in the profiles

→ genetic associations to <u>ANY</u> disease







## Main findings

Companies tested 69 polymorphisms in 56 genes. We found:

- 24/56 genes: no meta-analyses
- 7/56 genes: no statistically significant association to ANY disease
- 25/56 genes were associated with 28 different diseases
- Minor effects, most ORs 0.75-1.5; few rare exceptions
- Cardiogenomic profiles: genes were more frequently associated with non-cardiovascular diseases than with vascular diseases
- Osteogenomic profiles: 2 of 5 genes were associated with disease. Yet, NOT with bone diseases, but with Alzheimer disease, asthma, non-Hodgkin's lymphoma, obesity, psoriasis, SLE

Janssens et al. Am J Hum Genet 2008





### **VIEWPOINT**

## Uninformed consent in nutrigenomic research

A Cecile JW Janssens\*,1,2, Eline M Bunnik³, Wylie Burke⁴ and Maartje HN Schermer³

Informing also includes impact of tested genes on other diseases

Commentary reports about two studies

- Testing APOE gene to tailor recommendations saturated fat intake
- Forgot to disclose increased risk of Alzheimer's disease

Related ethical issues around informed consent, privacy, and data sharing



OMICS A Journal of Integrative Biology Volume 19, Number 9, 2015 Mary Ann Liebert, Inc. DOI: 10.1089/omi.2015.0109

### **Original Articles**

Meta-Analysis of Genes in Commercially Available Nutrigenomic Tests Denotes Lack of Association with Dietary Intake and Nutrient-Related Pathologies

Cristiana Pavlidis, Zoi Lanara, Angeliki Balasopoulou, Jean-Christophe Nebel, Theodora Katsila, and George P. Patrinos

### Methods

A thorough search was conducted in the PubMed literature database for genotype-phenotype correlation studies on 38 genes using the following terms: "nutrigenomics," "gene name," and "disease name" (Table 1). The 38 genes of interest were chosen following research on commercially available nutrigenomics tests and their specifics. Our metanalysis included 1170 entries published from 1995 to 2012.

Table 1. Search Criteria for Gene and Disease/Pathological Condition Association		
Genes	Disease/Pathological condition: investigation of possible association	
APOA1, APOA5,	Cardiovascular disease, Coronary heart disease, Coronary artery disease,	
APOB, APOC3, APOE	Hypercholesterolemia	
CETP	Cardiovascular disease, Coronary heart disease, Coronary artery disease, Hypercholesterolemia, Diabetes	
GJA4 (CX37)	Atherosclerosis, Hypertension, Stroke, Coronary artery disease	
HMGCR	Hepatic disease, Non- alcoholic fatty liver	
LIPC	Hepatic disease, Hypercholesterolemia, Fatty acid metabolism	
LPL	Dyslipidemia, CVD, Metabolic syndrome	
PON1	Atherosclerosis, Diabetes, Metabolic syndrome	
CAT	Diabetes, Hepatic diseases, Kidney diseases	
GPX1	Various types of gastrointestinal cancer	
GSTM1	Various types of gastrointestinal cancer, Various types of gastrointestinal disease, Coronary artery disease	
GSTP1	Various types of gastrointestinal cancer, Various types of gastrointestinal disease, Coronary artery disease	
GSTT1	Non-alcoholic fatty liver disease, Various types of gastrointestinal cancer, Various types of gastrointestinal disease, Coronary artery disease	
MNSOD	Various types of gastrointestinal cancer, Coronary artery disease, Various types of gastrointestinal disease	

controls) in a total of 1,170 entries were obtained. Conflicting findings indicated that there was a great incompatibility regarding the associations (or their absence) identified. No specific—and statistically significant—association was identified for any of the 38 genes of interest. In those cases, where a weak association was demonstrated, evidence was based on a limited number of studies. As solid scientific evidence is currently lacking, commercially available nutrigenomics tests cannot be presently recommended. Notwithstanding, the need for a thorough and continuous nutrigenomics research is evident as it is a highly promising tool towards

## What professionals say

right. 2014

FROM THE ACADEMY

**Position Paper** 

Position of the Academy of Nutrition and Dietetics: Nutritional Genomics

Genes Nutr (2013) 8:373-381 DOI 10.1007/s12263-013-0338-6

2013

REVIEW

Do we know enough? A scientific and ethical analysis of the basis for genetic-based personalized nutrition

Ulf Görman · John C. Mathers · Keith A. Grimaldi · Jennie Ahlgren · Karin Nordström

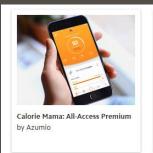
health messages to more individualized dietary guidance. Although the discipline of nutritional genomics holds promise for tailoring diet to a person's genotype and influencing chronic disease development, the science is still developing. The knowledge gained from nutritional genomics requires an evidence-based approach to validate

### Conclusion

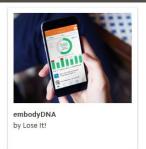
There is convincing evidence that common diet-related diseases are influenced by genetic factors, but knowledge in this area is fragmentary and few relationships have been tested for causality. The evidence that genotype-based dietary advice will motivate appropriate behavior changes is also limited. However, traditional nutritional advice is not always based upon causality but also on observational epidemiological studies. In several specific cases of genediet interaction, it may be more beneficial for identifiable groups of individuals with specific genotypes to follow personalized nutritional advice rather than general dietary recommendations. From an ethical perspective, a precau-



Now you can eat a healthy diet best suited to your genetic makeup, your metabolism, and your lifestyle. Turn DNA insights into easy-to-make food decisions for balanced meals that make your body happy.







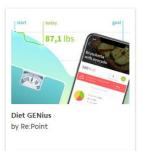






Heart Optimizer: Genetics and Nutrition Coaching by Arivale







### Nutrition

## Calorie Mama: All-Access Premium by Azumio

Lose weight the smart way with nutrition and workout plans customized to your DNA. Easily count calories by taking photos of your meals. Available exclusively on iOS.





## What companies promise



Sign In



DNAFit UNLOCKYOUR HUMAN POTENTIAL

Now it's your turn - this is how it works

You're not the same as everybody else, so why follow everybody else' diet?









Gem right

## Take the guesswork out of dieting with a personalized nutrition plan based on your DNA





Whether you're looking to slim down, maintain, or just want to eat a little healthier, your genetics contains valuable information to take control of your health and wellness in ways never before possible.

It's simple — your body is unique and your weight loss journey should be as well. Your DNA plays a large role in determining how your body interacts with food — from your preferences to your sensitivities and metabolism. This means that no generic diet is right for everyone.



People on a genetic-based diet lost 33% more weight on average than those on regular diet plans.

ep ng, and Sign In

lix, you ess to a DNA-oducts Ip you formed e longer, and le best hly need DNA once.



## What their disclaimers reveal

MealPlanner

by DNAFit

Overview

The science

Ouestions

∰ Gift

Order

### The science

Genetic variants related to nutrition are connected to the way that your body processes food, but they do not guarantee that you will or will not be successful with any given diet plan. Your DNA may help you narrow in on new diet plans that you might prefer or find more successful than others, or even just a better understanding of your existing preferences. Everyone, regardless of their genetics, will benefit from a well-balanced diet.

#### **Impact**



Mostly other factors

Genetics can help, but your diet has the biggest impact on your nutrition.

#### Limitations

- This product will not provide any medical information.
- Genetic results are based on population-wide studies. What is true at a population level average may not be true at an individual level.
- Results may be more accurate for individuals of certain genetic backgrounds.
- Results do not determine or limit your ability to gain or lose weight.
- Our understanding of how genetics influences metabolism will improve with more research.



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## **Ethical principles**

### Medicine

- Autonomy
- Beneficence
- Nonmaleficence
- Justice

### Marketing (AMA)

### Ethical norms:

- Do no harm
- Foster trust in marketing
- Embrace ethical values:
   Respect, Honesty,
   Responsibility,
   Transparency, Fairness,
   Citizenship

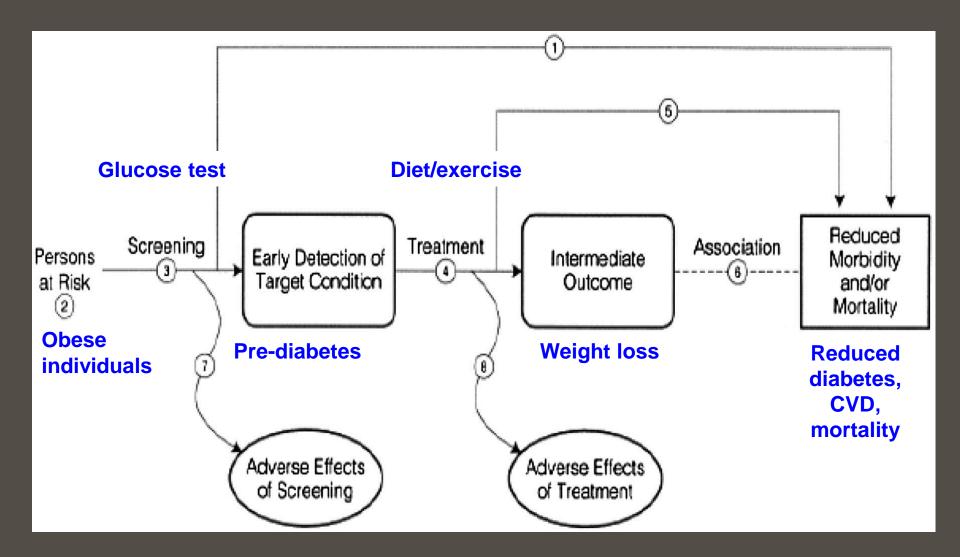


## Beneficence

- Intent of doing good
- Developing and maintaining skills and knowledge, and continually update
- Consider individual circumstances of all patients
- Is the commercial offer in the interest of customers or the company?
- Do we have evidence on how to 'compensate' genetic effects with diet? (no)

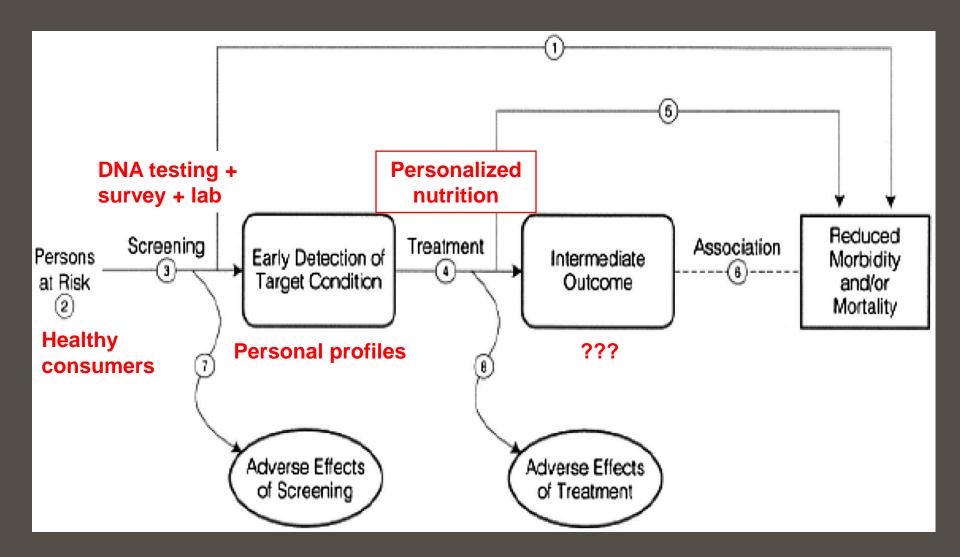


## **USPSTF Analytic Framework**





## **USPSTF Analytic Framework**





## **Autonomy**

 Customer should decide whether testing is useful for them → personal utility

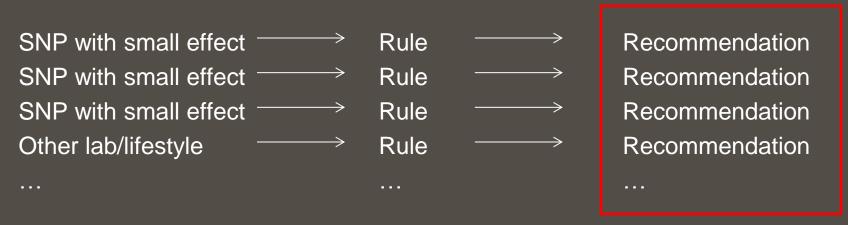
- Needs information to make informed decision making
- Companies use proprietary algorithms give no insight in validity of recommendations

Customers and scientists cannot verify validity



## Proprietary algorithms: how advanced?





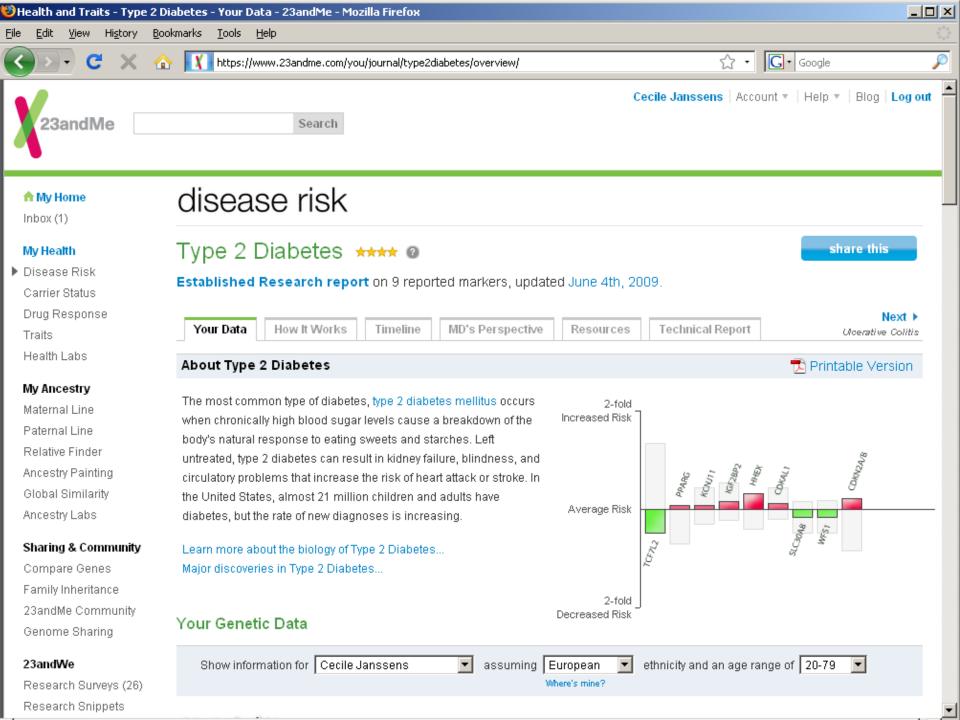
Personalized combination



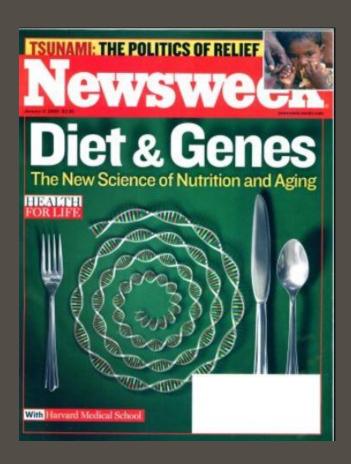
Table 3: Personalized recommendations given to the Nutrigenetic test patient group in addition to base diet.

Nutrient intervention group	% Receiving modified advice
Variation in MTHFR, MTRR, MTR or CBS:	98.6
Rationale: Polymorphisms in genes involved in folic acid metabolism have been shown to influence this pathway affecting plasma homocysteine levels as well as the balance between DNA methylation and synthesis of nucleotides [14, 15].	
Recommendation: Add supplement containing 800 mcg folic acid, 15 mg Vitamin B6 and 20 mcg B12	
Variation in GSTM1, GSTT1 or GSTP1:	76.1
Rationale: Patients with deletions in GSTM1 which affect Phase II detoxification processes have been shown to have reduced levels of DNA adducts [16], and increased levels of GSTA1 circulating activity [17], when adequate levels of cruciferous vegetables have been consumed. Risk for lung cancer drops by up to 80% in individuals lacking GSTM1 and/or GSTT1 genes when consumption of cruciferous vegetables is high [18].	
Recommendation: Ensure diet includes regular portions of cruciferous (5 times per week) and allium (daily)	
vegetables (suggestions and recipes provided to patient). Add broccoli extract and allium supplement if required. Variation in SOD2, SOD3, NOS3:	48.6
Rationale: superoxide dismutase enzymes are free radical scavengers that have important antioxidant activity which can be affected by genetic polymorphism [19]	
Recommendation: Add supplements containing antioxidants, Vit A (5,000 IU), Vit C (250 mg) and Vit E (200 IU). Variation in VDR, COLIAI:	87.5
Rationale: Several studies have shown that gene-diet interactions have a role to play in maintenance of bone condition. For example caffeine increased rate of bone loss but only in the presence of the VDR taq1 variant [20]. Others have shown gene-diet effects involving calcium [21, 22] and vitamin D [23].	
Recommendation: Keep caffeine below 2 cups coffee/day. Increase dairy component of diet (yoghurt, cheese and low fat milk). If required add supplement containing 800 IU vitamin D and 1,300 mg Calcium	
Variation in TNF $\alpha$ , IL6, NOS3: Rationale: Variations in inflammation pathway genes TNF $\alpha$ and IL6 have been shown to be pro-inflammatory and the effect can be modulated by increased levels of fish oil in the diet [24]	65.3
Recommendation: Add supplement Omega 3 (700 – 1,400 mg). Make sure weekly diet contains portions of oily fish	
Variation in CETP, LPL, APOC3:	79.2
Rationale: Polymorphisms in genes involved in lipid metabolism and transport, in combination with dietary fat intake, have been shown to affect plasma cholesterol levels [25]	
Recommendation: The base low fat is already within the limits recommended for these variations so no further specific advice is given but current advice is reinforced and advice given to restrict consumption of dairy foods.	
Variation in ACE, PPARG:	80.6
Rationale: gene-diet and gene-exercise interactions have been reported to affect blood glucose and insulin levels [26, 27]	
Recommendation: The base low glycemic diet is already within the limits recommended for these variations so no further specific advice is given but current advice is reinforced. Extra exercise advised for this group	





### **Onward**



- Better and more relevant scientific studies
- More respectful conversation with consumers
  - Genetically personalized
     nutrition recommendations is still
     premature, largely lacking
     appropriate scientific basis
  - Just say it

2005

