What is Precision Public Health and Where Does FH Fit in?

Hot Topics of the Day

Weekly Update

Muin J. Khoury, MD, PhD

Office of Genomics and Precision Public Health Centers for Disease Control and Prevention



Family Health History

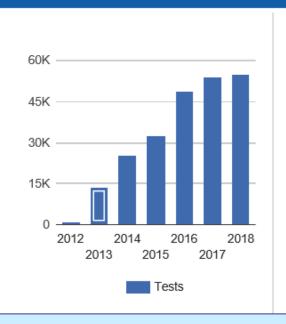
Genetic Counseling & Testing



My Family Health Portrai

From Genomic Medicine to Precision Medicine

- Increasing Number of genetic tests (from single genes to panels and NGS)
- Whole-genome sequencing as tool in clinical and public health practice
- Increasing public interest and proliferation of direct-to-consumer tests
- All of Us Research Initiative (1M person cohort)- genes and environments



What is precision medicine?

"An emerging approach for disease prevention and treatment that takes into account people's individual variations in genes, environment, and lifestyle."

CDC Office of Genomics and Precision Public Health

1. Identifyevidence-basedapplications

2. Inform and communicate

3. Integrate into practice & programs

- Selected "Tier 1" applications
- BRCA1/2 Related Cancers, Lynch Syndrome
- Familial Hypercholesterolemia
- Collectively Affect >2 Million People in US and Most Don't know it
- Implementation of existing evidence-based guidelines can prevent cancer & heart disease, & save thousands of lives every year!
- Toolkit for public health departments
- Working with CDC programs and external partners

Will Precision Medicine Improve Population Health?



Will Precision Medicine Improve Population

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Sandro Galea, MD,

Boston University School of Public Health Boston, Massachusetts Health?

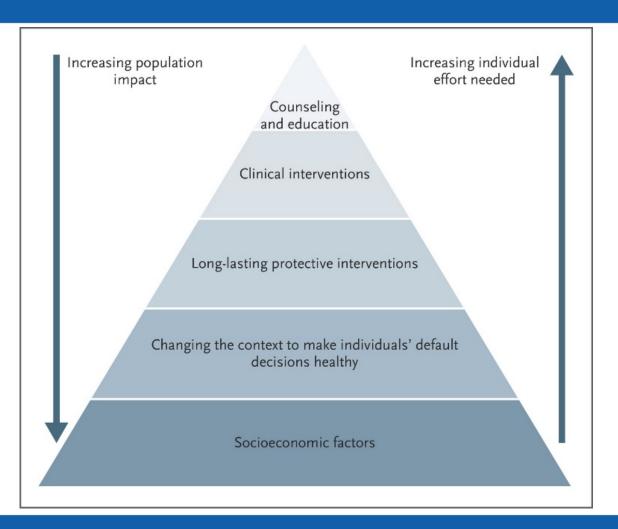
Announcement of the precision medicine initiative has led to a variety of responses, ranging from enthusiastic expectations¹ to explicit skepticism,² about potential health benefits, limitations, and return on investment. This Viewpoint discusses whether precision medicine is unlikely or likely to improve population health, aiming to forge a consensus that bridges disparate perspectives on the issue. The potential of precision medicine to improve the health of individuals or small groups of individuals is not addressed here because it involves a different question with different metrics.

Precision Medicine Is Unlikely to Improve **Population Health**

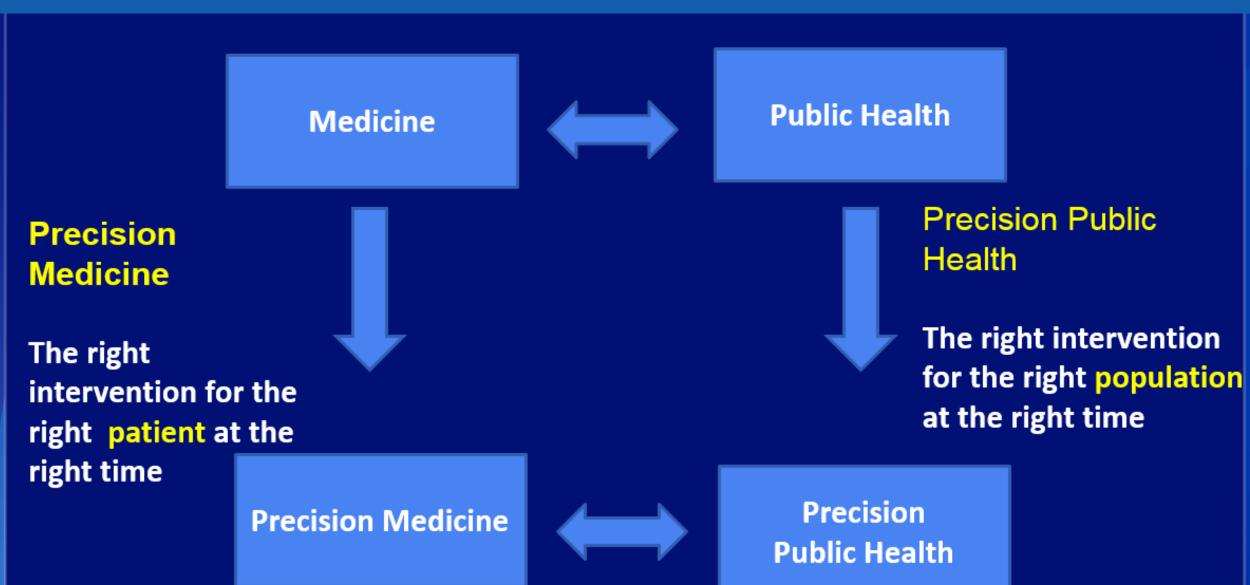
There are 3 fundamental reasons that precision medicine might not improve the health of populations. First, disease pathogenesis, especially for common noncommunicable diseases, is extraordinarily complex. Abundant evidence has demonstrated this for the association between the multiplicity of specific genes and conditions, including obesity, hypertension, or certain cancers. Additionally, it is known that genetic associa-

health of its population. Over the past 30 years, the United States has fallen behind other high-income peer nations in health attainment on many metrics, including life expectancy and infant mortality, and there are persistent gaps in health outcomes by income and race/ethnicity. 4 The solution to these challenges is probably not an increased focus on the individual, but rather lies in focusing on the social, economic, and structural drivers of population health that are ubiquitous and inevitably linked to health achievement as a country. The centrality of the precision medicine effort to the US national health research agenda may distract from efforts to remedy the foundational causes of ill health such as poverty, obesity, and education. Without addressing these causes, there will be little, if any, success in reversing the trends of poor achievement in US population health.

Second, precision medicine could (and to some extent has) led to a shift from which projects are funded by health research agencies. Funding for grants with a population health or public health goal has declined over the past 10 years at the National Institutes of Health.



Precision Medicine and Precision Public Health: Two Peas in a Pod!



Public Health Is Increasingly Driven by Big Data!

Beyond Public Health Genomics: Can Big Data and Predictive Analytics Deliver Precision Public Health?

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PHG, 2019

Keywords

Big data · Genomics · Implementation science · Medicine · Predictive analytics · Public health

Abstract

The field of public health genomics has matured in the past two decades and is beginning to deliver genomic-based interventions for health and health care. In the past few years, the terms precision medicine and precision public health have been used to include information from multiple fields measuring biomarkers as well as environmental and other variables to provide tailored interventions. In the context of public health, "precision" implies delivering the right intervention to the right population at the right time, with the goal of improving health for all. In addition to genomics, precision public health can be driven by "big data" as identified by volume, variety, and variability in biomedical, sociodemographic, environmental, geographic, and other information. Most current big data applications in health are in elucidating pathobiology and tailored drug discovery. We explore how big data and predictive analytics can contribute to precision public health by improving public health surveillance and assessment, and efforts to promote uptake of evidence-

based interventions, by including more extensive information related to place, person, and time. We use selected examples drawn from child health, cardiovascular disease, and cancer to illustrate the promises of precision public health, as well as current methodologic and analytic challenges to big data to fulfill these promises. 6 2019 S. Karger A.G. Basel

Introduction

The term "precision" is increasingly used in medicine [1] and public health [2]. Precision medicine is often used synonymously with genomic medicine, and precision public health has been equated with applications of precision medicine in populations [3]. Applications of precision medicine (e.g., cancer genomics) are unlikely to lead to improved population health, as targeted interventions benefit only a small subset of the population. Nevertheless, as we and others have discussed, there is a bigger role for precision in public health beyond genomics [4]. Precision public health can be viewed as the delivery of the right intervention to the right population at the right time, and includes consideration of social and environ-

Put at its simplest, precision public health is about using the best available data to target more effectively and efficiently interventions of all kinds to those most in need."

R Horton, Lancet 2018

Can we Conduct Public Health Functions With More "Precision"?

Assessment

- More "precision" in measuring population health problems
- Policy Development
 - Developing the right intervention for the right population at the right time
- Assurance
 - More "precision" in delivering interventions & addressing health disparities

Precision Public Health for the Era of Precision Medicine

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he Precision Medicine Initiative promises a new healthcare era. A proposed 1 million-person cohort could create a deeper understanding of disease causation. Improvements in quality of sequencing, reduction in price, and advances in "omic" fields and biotechnology promise a new era, variably labeled personalized or precision medicine. Although genomics is one driver of precision health care, other factors may be as important (e.g., health information technology).

Both excitement and skepticism met the announcement.² Public health experts are concerned about the disproportionate emphasis on genes, drugs, and disease, while neglecting strategies to address social determinants evidentiary foundation for use. The following are examples of priority areas.

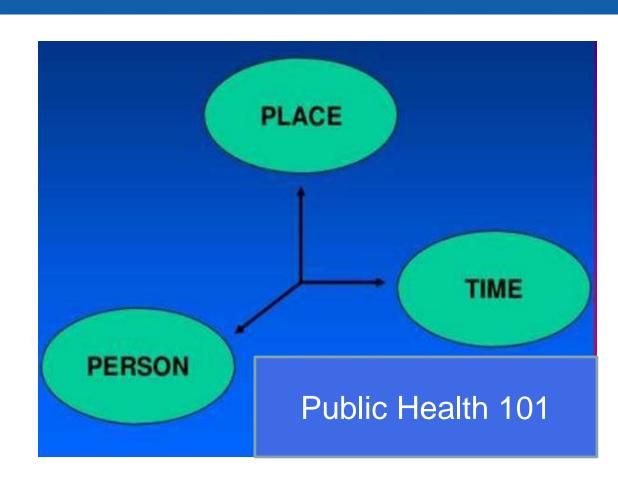
AJPM, 2016

Role of Multidisciplinary Public Health Sciences

Though precision medicine focuses on individualized care, its success truly requires a population-based approach. To learn what interventions work for whom, data on each individual need to be compared with data from large, diverse numbers of people to identify population subgroups likely to respond differently to interventions. In addition, collecting information from

Using Big Data in Public Health Requires Complex Analytics, By Person, Place and Time

- Person
 - Disease subtypes using biomarkers
 - Disease susceptibility (e.g. genetics)
- Place
 - Geographic and environmental data
 - > Health systems
- ☐ Time
 - Wearables
 - Social Media



Khoury MJ et al, Public Health Genomics, 2019

Examples of Precision Public Health: 1) Influenza Surveillance, 2) "Spotting Obesity from Space"

ARTICLE

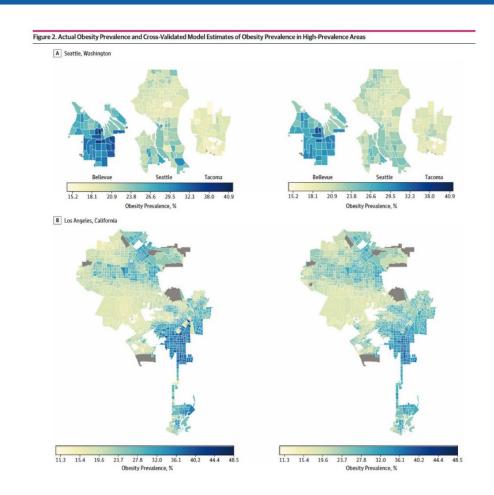
https://doi.org/10.1038/s41467-018-08082-0

OPEN

Improved state-level influenza nowcasting in the United States leveraging Internet-based data and network approaches

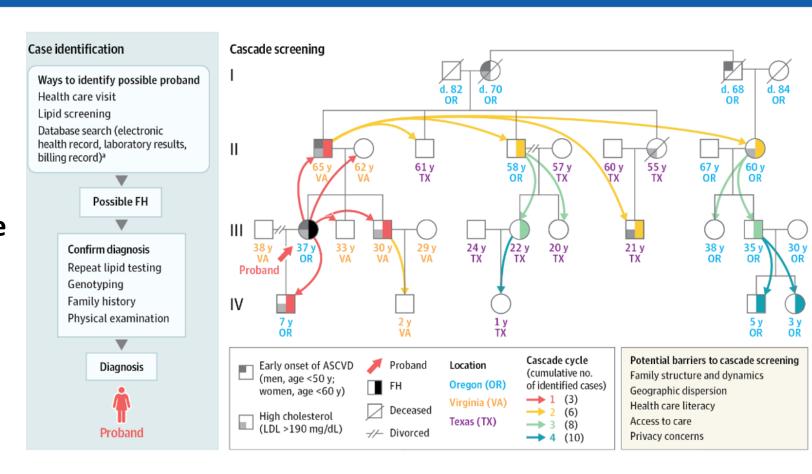
Fred S. Luo 1, Mohammad W. Hattab², Cesar Leonardo Clemente³, Matthew Biggerstaff⁴ & Mauricio Santillana^{1,5}

In the presence of health threats, precision public health approaches aim to provide targeted, timely, and population-specific interventions. Accurate surveillance methodologies that can estimate infectious disease activity ahead of official healthcare-based reports, at relevant spatial resolutions, are important for achieving this goal. Here we introduce a methodological framework which dynamically combines two distinct influenza tracking techniques, using an ensemble machine learning approach, to achieve improved state-level influenza activity estimates in the United States. The two predictive techniques behind the ensemble utilize (1) a self-correcting statistical method combining influenza-related Google search frequencies, information from electronic health records, and historical flu trends within each state, and (2) a network-based approach leveraging spatio-temporal synchronicities observed in historical influenza activity across states. The ensemble considerably outperforms each component method in addition to previously proposed state-specific methods for influenza tracking, with higher correlations and lower prediction errors.



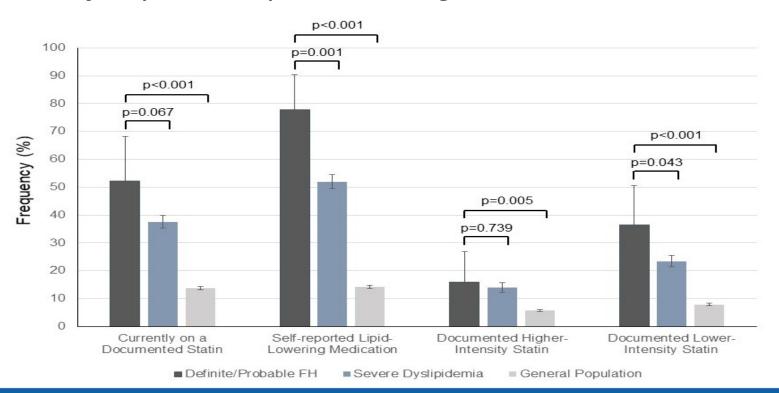
Familial Hypercholesterolemia

- Common autosomal dominant condition associated with premature death from heart disease (> 1M affected in US)
- Evidence-based recommendation for aggressive cholesterol reduction and cascade screening in relatives
- Highly underdiagnosed and undertreated
- Racial and ethnic disparities in diagnosis and management



Familial Hypercholesterolemia is Common and Undertreated in the United States

Prevalence of documented statin and self-reported lipid lowering medication use



Young and uninsured patients are at the highest risk for under treatment

Familial Hypercholesterolemia in the Age of **Precision Public Health**

RESEARCH LETTER

Use of Twitter to Promote Awareness of Familial Hypercholesterolemia

wareness of familial hypercholesterolemia (FH), a prevalent genetic disorder that greatly increases risk of early onset myocardial infarction, is low." It is estimated that of the =1.3 million people with FH in the United States, <10% have been diagnosed.1 Twitter is being increasingly used in academic cardiovascular settings?; however, its potential for promoting awareness of cardiovascular genetic disorders such as FH is unknown. September is designated as the National Cholesterol Education Month (NCEM) to promote awareness of hypercholesterolemia as a risk factor, including through social media. In addition, the International FH Awareness Day established by the FH Foundation as part of NCEM, includes a Tweet-a-thon. We investigated whether there was an increase in FH-related Twitter activity during NCEM in September 2018, using this as a surrogate for FH awareness.

This study was deemed exempt by the institutional review board as it did not involve human or animal subjects. Publicly available Twitter data were used to support the findings of this report and are available from the corresponding author

We quantified FH-related Twitter activity from August 2018 to October 2018 using Sprinklr Listening Explorer software to extract tweets and engagement metrics, including reach (the potential audience size of tweets, measured by total follower counts of users who shared content). As a control, we examined the reach of colorectal cancer-related tweets during the same period.

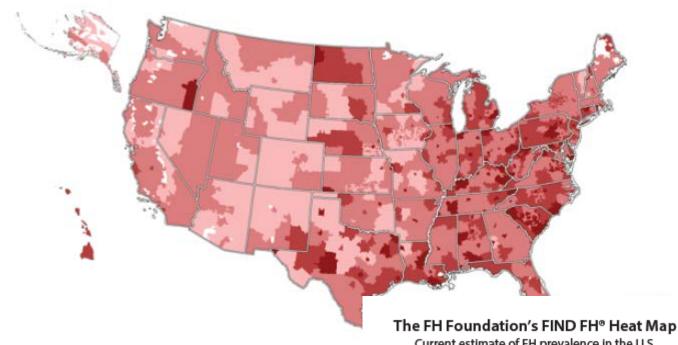
FH-related tweets were classified as true positive or false positive based on a keyword list derived from manual review of tweets. This preprocessing step was performed by keyword matching using Python (script is publicly available and can be accessed at https://github.com/bastia1989/FHtweet). All true positive tweets were then categorized via name entity recognition of profile descriptions, to identify users as organizations or individuals. Tweets were also classified based on their message format as updates (mentions+original tweets+retweets with comments), retweets (without comments), and replies.

A qualitative content analysis of a random sample of 600 tweets was performed, and topic-keyword tables were created to define 6 following thematic categories: awareness/prevention, diagnosis, treatment, lifestyle, knowledge dissemination, and conferences/fundraisers. Tweets were then binned into these categories based on presence of relevant keywords.

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FAMILIAL HYPERCHOLESTEROLEMIA

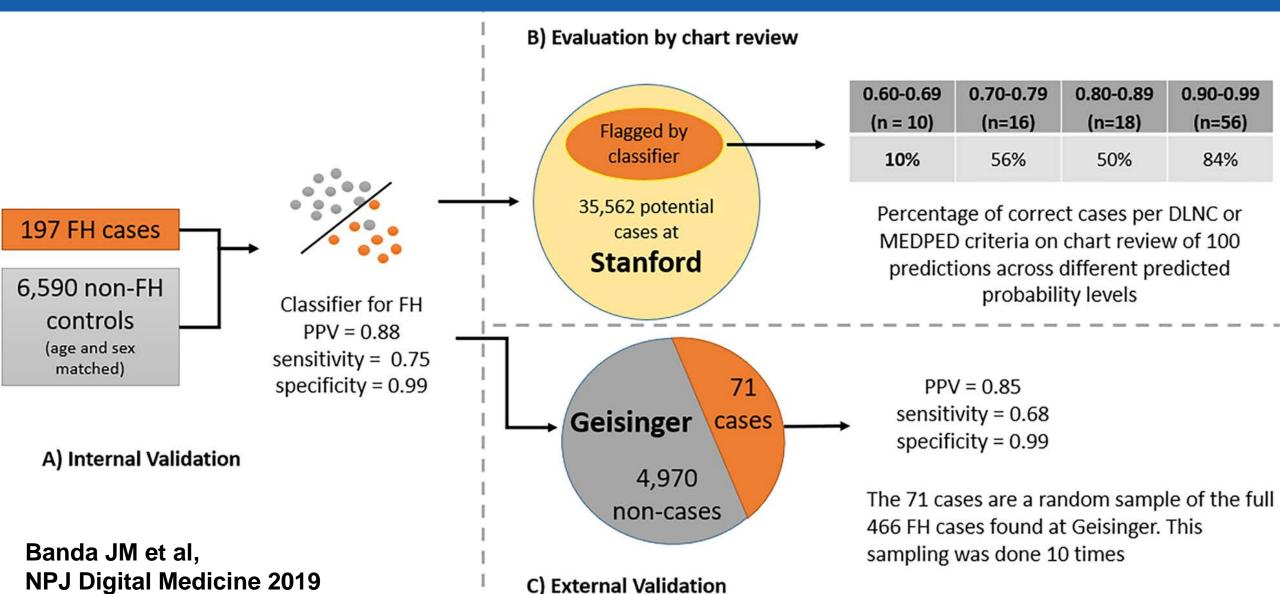
The most common cause of early and agressive heart disease



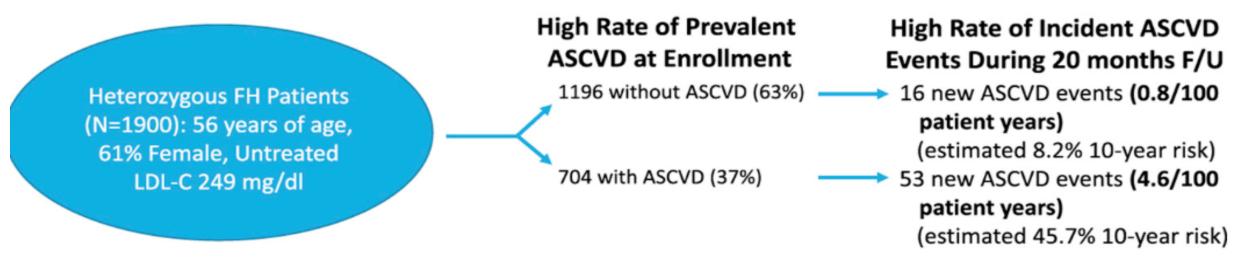
Current estimate of FH prevalence in the U.S. Estimated prevalence of FH with > 0.66 probability by zip code

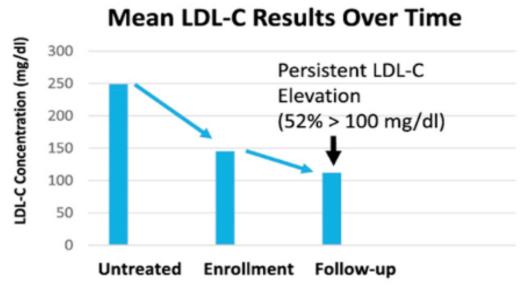
> < 0.20% – Less than 1 in 500 0.20% to 0.39% - 1 in 500 to 1 in 250 0.40% to 0.59% - 1 in 250 to 1 in 165 ≥ 0.60% - Greater than 1 in 165

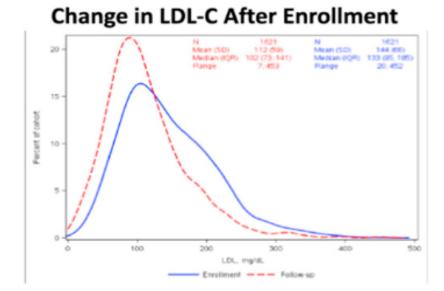
Finding Missed Cases of Familial Hypercholesterolemia in Health Systems Using Machine Learning



CASCADE FH REGISTRY Longitudinal Follow-Up Analysis







PB Duell et al.. Atherosclerosis 2019

FH in the Era of Precision Public Health

