



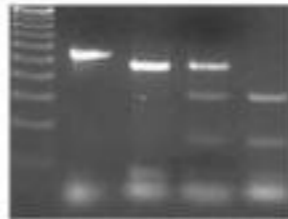
P6 – GENOMICS TO HEALTHCARE

Prevention of common chronic diseases using modern medicine



The march of technology...

single variant
(10^0 SNPs; 10^3 genotypes)



detailed study of individual genes
(10^2 SNPs; 10^{5+} genotypes)



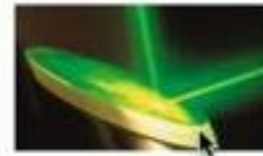
regional studies
(10^4 SNPs; 10^8 genotypes)



genome-wide association
(10^6 SNPs; 10^{10} genotypes)

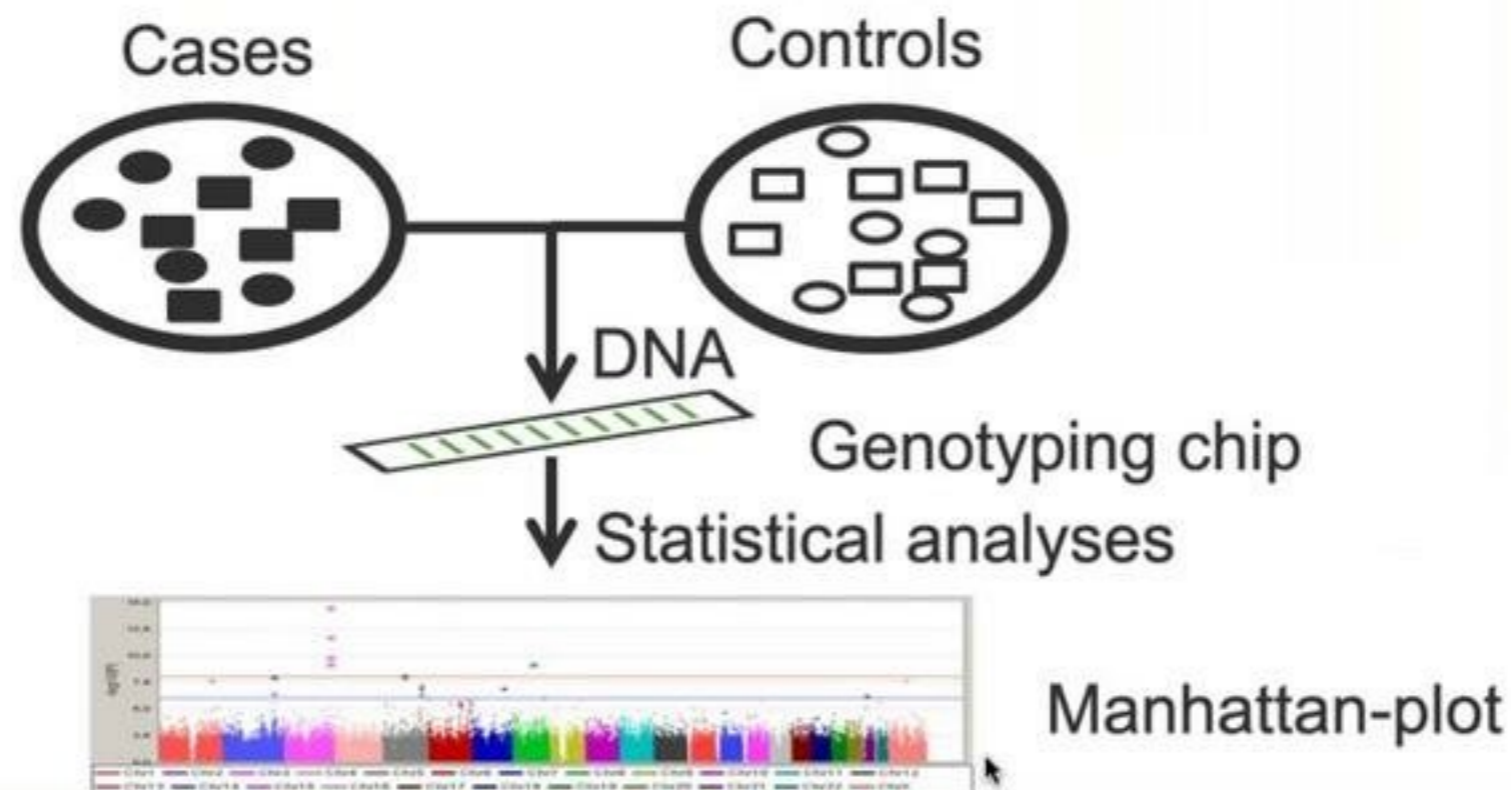


complete resequencing
(10^8 SNPs / 10^{12} genotypes)



WHAT IS A GWAS?

- Genetic (chip) information is used to evaluate whether the genetic marker(s) are associated with a phenotype (qualitative or quantitative)





NHGRI-EBI GWAS Catalog
www.genome.gov/GWASStudies
www.ebi.ac.uk/fgpt/gwas/

CORONARY HEART DISEASE GWAS



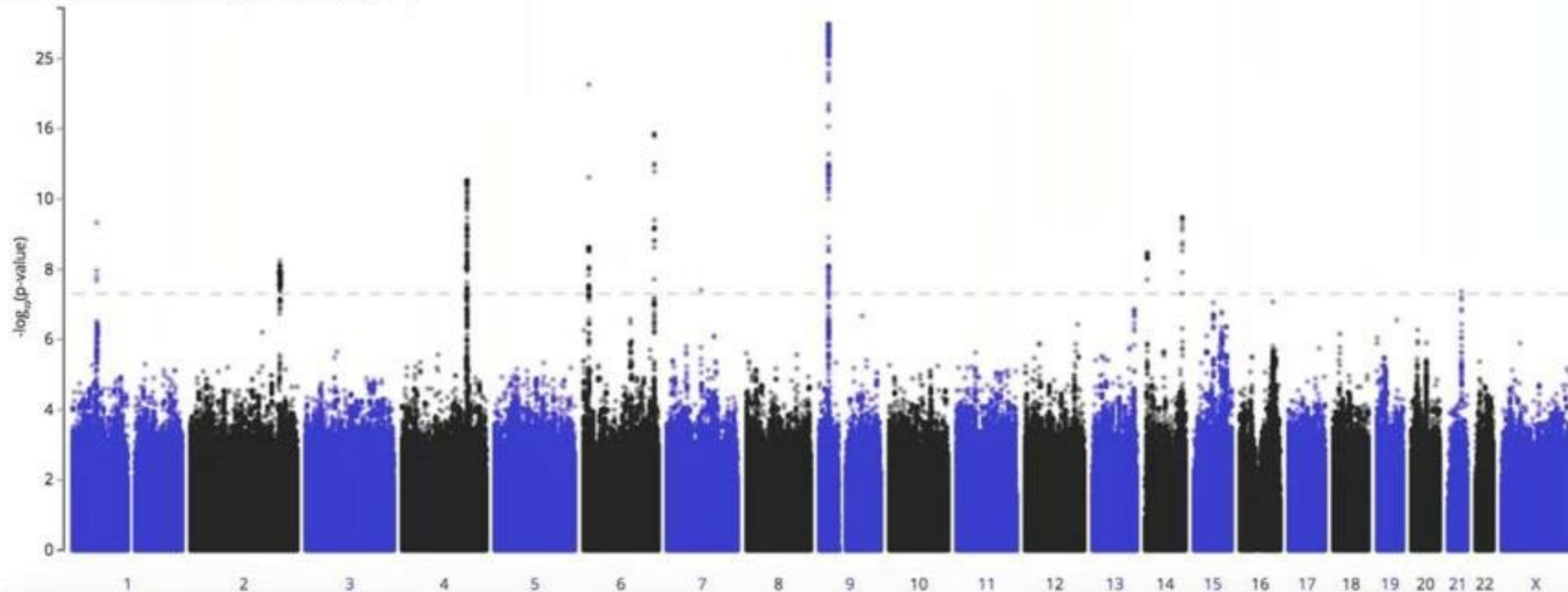
Major coronary heart disease event

Category: Circulatory system

FINN GEN: 13753 cases, 121885 controls

UKBB: 10157 cases, 351037 controls

p-values smaller than $1e-10$ are shown on a log-log scale



PRS = A NOVEL TOOL TO ESTIMATE GENOMIC RISK

KP Yhteiskäsittely...
<https://tila.tilmeri.fi/User%20Guide/>

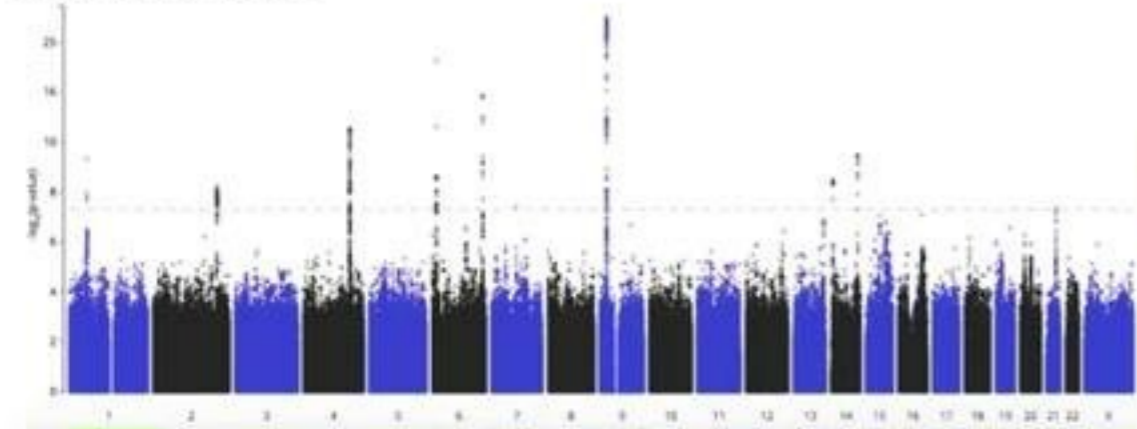
Vastaus

- Most common diseases are multifactorial and polygenic
- GWAS data can be used to form so-called Polygenic Risk Scores (PRS) which condense the risk produced by her/his genomic variation of an individual to a single number for a given phenotype.

Major coronary heart disease event

Category: Circulatory system

SNPs: 12758 cases, 127495 controls
Locus: 10157 cases, 351927 controls
Features smaller than 10 are shown on a log-log scale

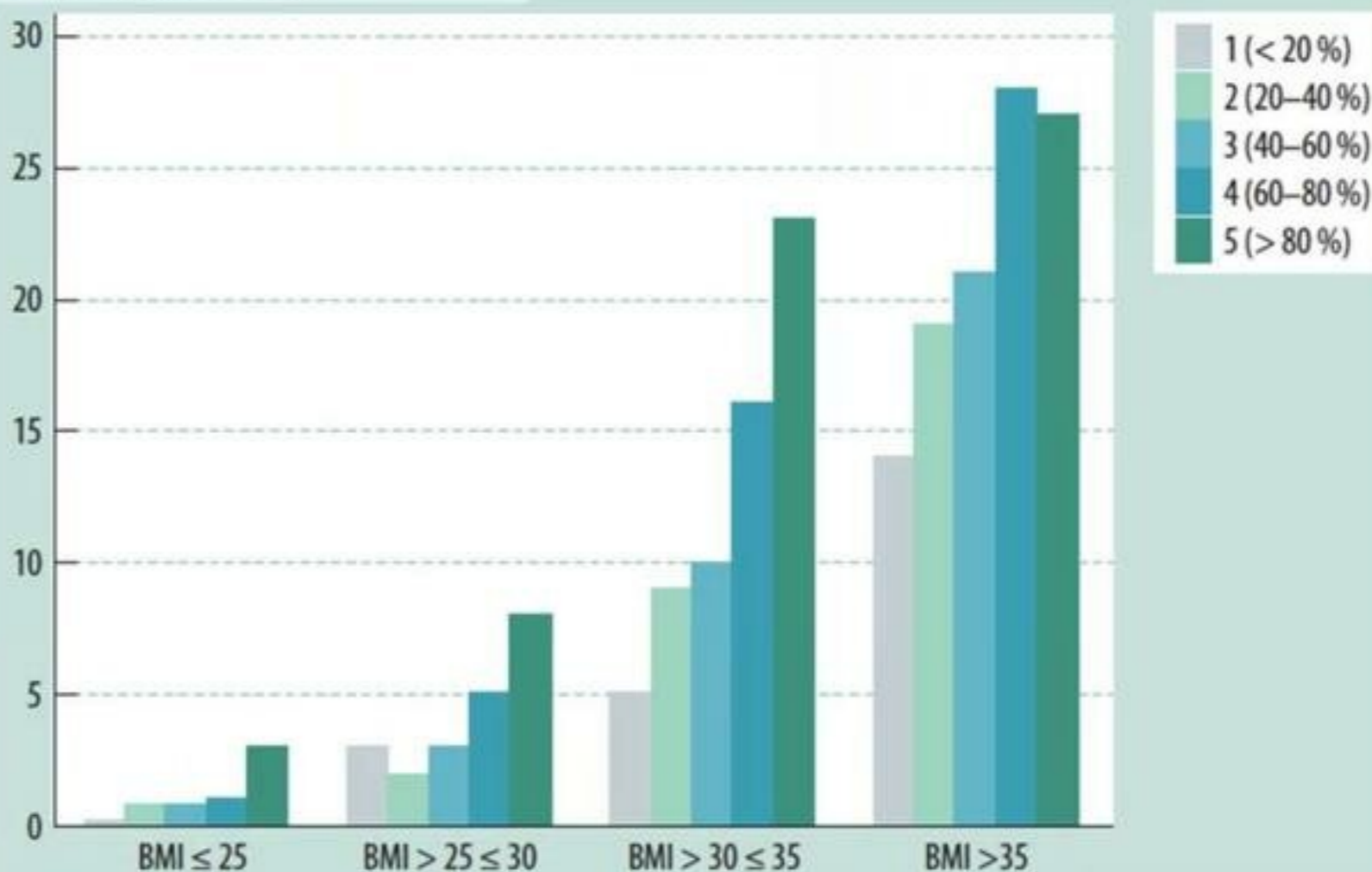


COMPARISON OF PRS AND BMI IN TYPE 2 DIABETES



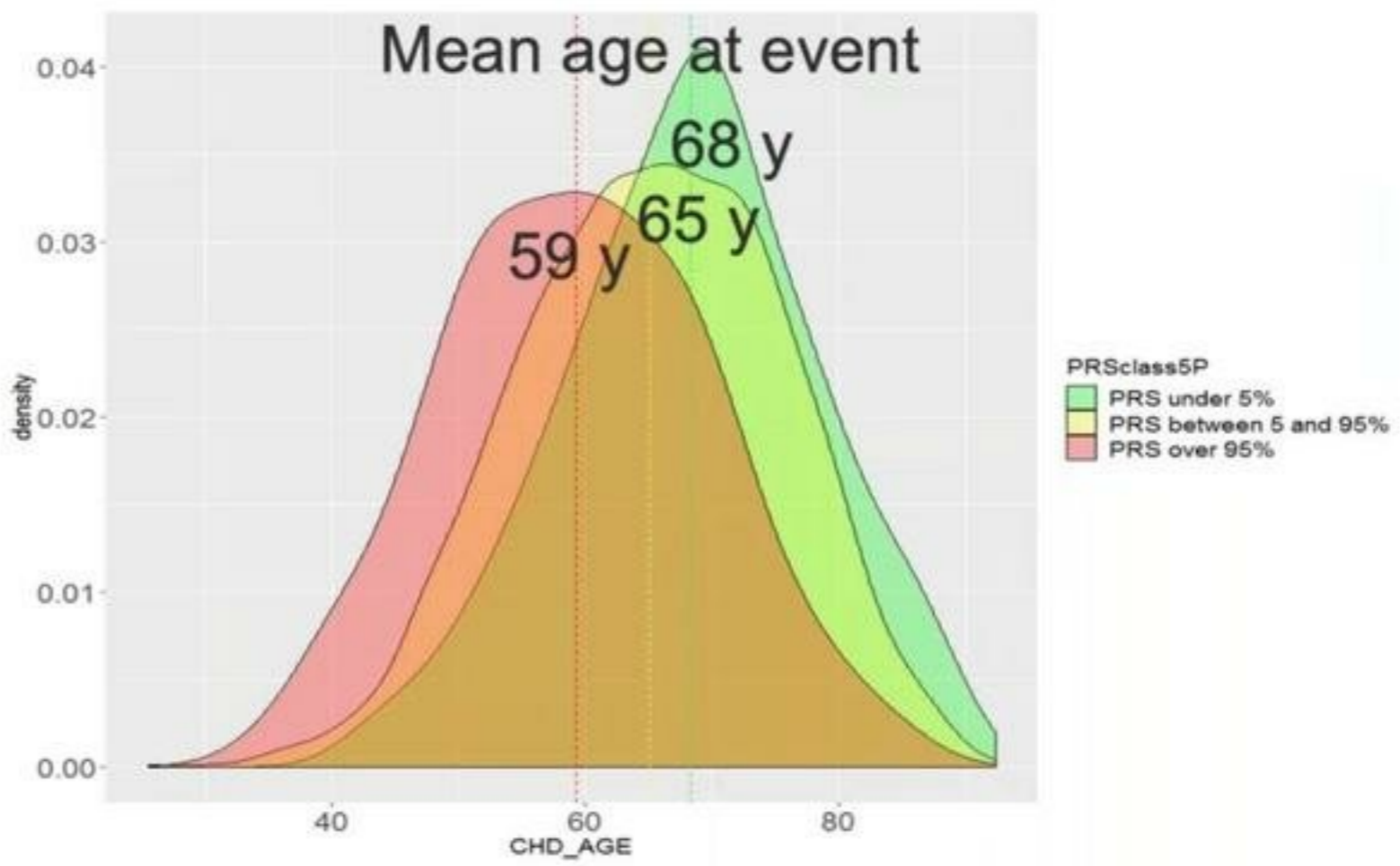
Genetic risk brings added personalized value to the disease prevention regardless of lifestyle

No (%) of type 2 diabetes cases in a 10-year follow-up



PRS ASSOCIATES WITH THE AGE OF THE FIRST CHD EVENT

High PRS may identify individuals who get a disease at early age



PRS IS NOT INSIGNIFICANT



But it is an experimental value, which is not yet accepted as a clinical tool



The impact of communicating genetic risks of disease on risk-reducing health behaviour: systematic review with meta-analysis

Gareth J Hollands,¹ David P French,² Simon J Griffin,³ A Toby Prevost,⁴ Stephen Sutton,³ Sarah King,¹ Theresa M Marteau¹

BMJ: first publi

“...communicating DNA based risk estimates changes behaviour is not supported by existing evidence ... do not support use of genetic testing or the search for risk-conferring gene variants for common complex diseases on the basis that they motivate risk-reducing behaviour.”

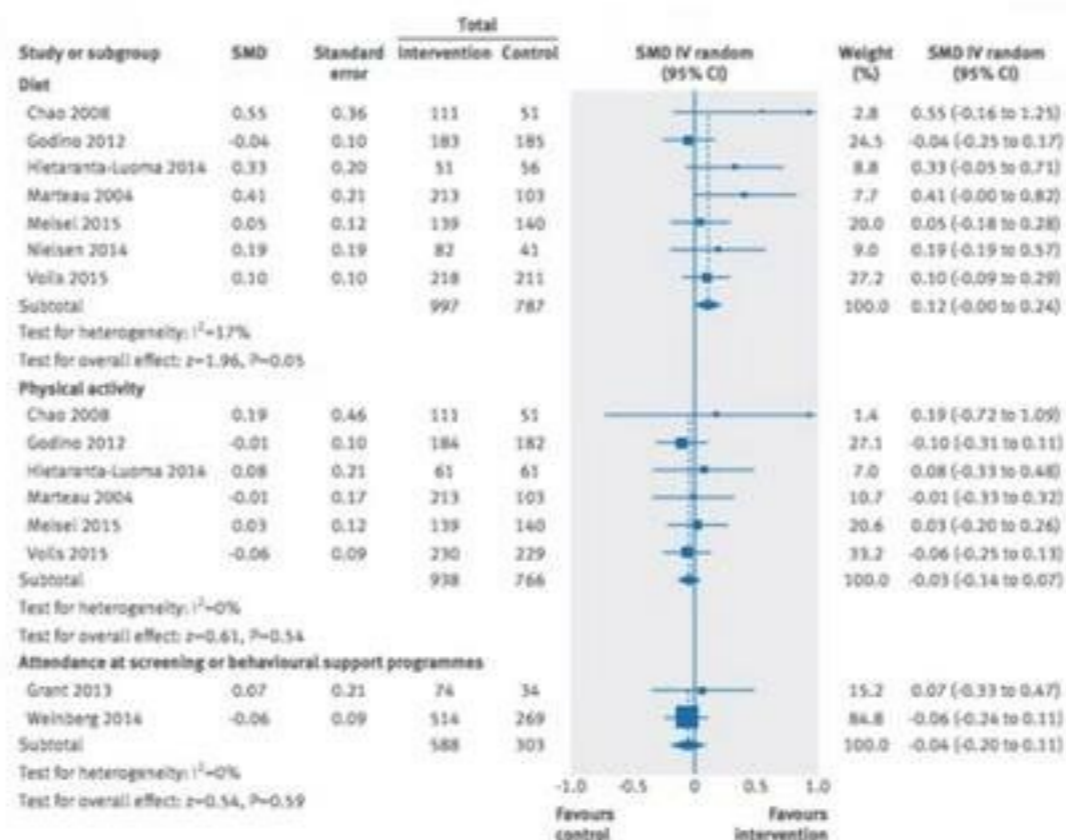


Fig 4 | Primary outcome analysis: diet; physical activity; attendance at screening or behavioural support programmes. SMD—standardised mean difference



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PUBLIC HEALTH PROGRAMMES

e.g. North Karelia Project 1972–, Finnish Allergy Program 2008–2018, Suicide Prevention Project, Indoor Air and Health

INTERVENTION STUDIES

e.g. Diabetes Prevention Study (DPS), FINGER, vaccine trials

HEALTHCARE

Focus heavily on treatment instead of prevention

Ageing population; increasing costs resulting from the chronic diseases

GENOMICS

A plethora of genomic regions found to associate with various diseases and traits (e.g. coronary heart disease, type 2 diabetes)

Costs of genome analyses rapidly decreasing

P6

POPULATION FOLLOW-UP STUDIES

e.g. HES-surveys: Mobile Clinic, FINRISK, Health 2000/2011, FinHealth

DATA RECORDS AND SAMPLE COLLECTIONS

e.g. national registries, biobanks

P6 – GENOMICS TO HEALTHCARE

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Objectives of the P6 initiative

Improved public health

Cost-effective use of the healthcare resources

Health equality of citizens

Finland as a healthcare pioneer

Improved prevention and early care of the common chronic diseases

Professionally directed use of genetic information in the clinical care

Novel opportunities for research, public-private co-operation and businesses in the health sector

Healthcare professionals have a new predictive, preventive, personalized and participatory tool for their clinical work that is ethically sound and simple to use

PURPOSE

Management and coordination

P6 Study

Communications

Health-economic evaluations

Public-private collaboration

Technology and biobanks

Training

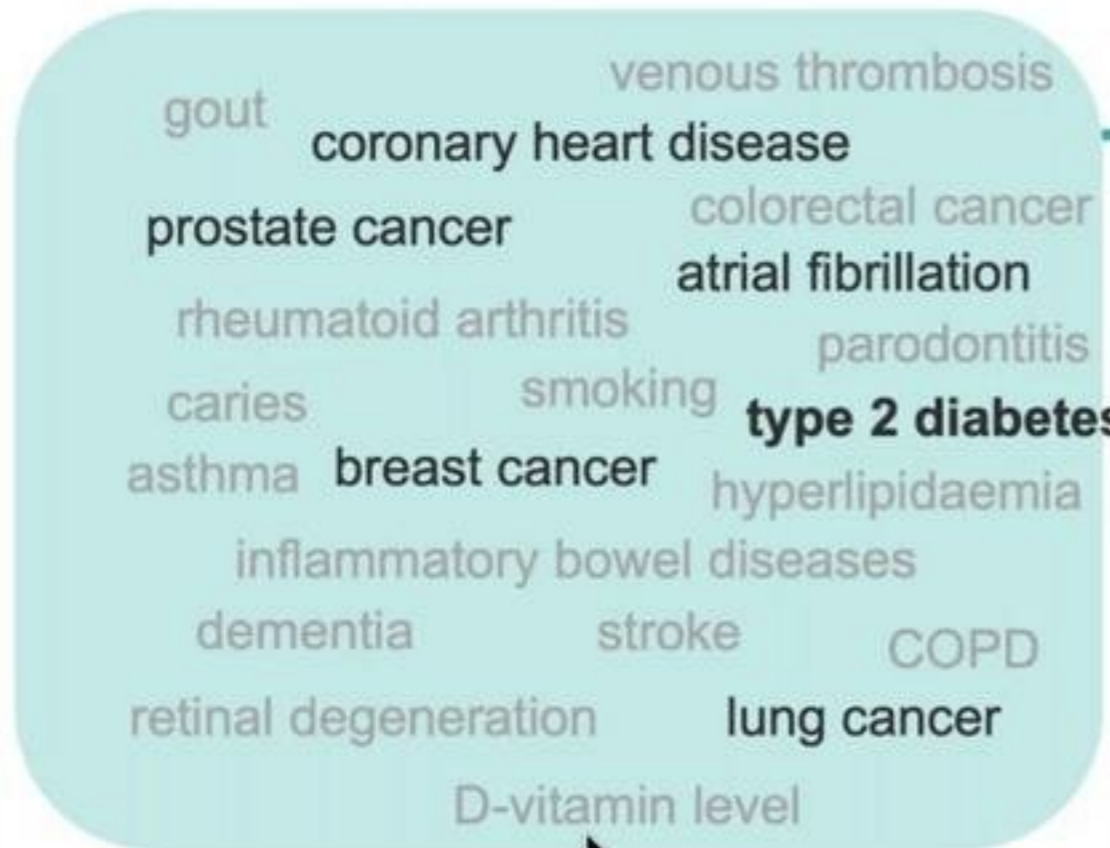


P6 – GENOMICS TO HEALTHCARE

Prevention of common chronic diseases using modern medicine



Disease endpoints selected to the P6 study



Forming polygenic risk scores (PRS) based on previous high-quality research. Evaluation of PRSs in a large prospective Finnish population sample collection

Recruitment of 100.000 participants from the Finnish biobanks. Returning disease risk information (genetic and lifestyle) via a secure web portal
900.000 registry controls

Invitation of 4.000 high-risk individuals to an active intervention (RCT)



P6 – TIMELINE



Strategic planning of the P6 initiative
2019

P5 FinHealth pilot
2018-2019

Recruitment of P6 study subjects via a biobank portal
2021-2022

Returning P6 risk information via a secure web portal
2022-2023

P6 PRS analyses in the THL cohorts
2020

P6 interventions
2021-2023

Analyses of the P6 results
2022-2024



Follow-up through national healthcare registries
Follow-up studies
2024 ->

- Kutsu kettin
- Saadatus ja kyseilylomake THL:een
- Riskilänsiden tarkastuksen alustamis keräilyssä suomalaisissa aivotietoisu ja genotyyppi
- Henkilökohtaisen riskilänsiden tarkastus
- Tulokset saatavilla erikseen OmapS-raportissa
- 26b
- Kysely OmaPS-sivustolla
- 2a
- Kysely OmaPS-sivustolla
- 2c
- Kysely OmaPS-sivustolla
- 2d
- Kysely OmaPS-sivustolla
- 2e
- Kysely OmaPS-sivustolla
- 2f
- Kysely OmaPS-sivustolla

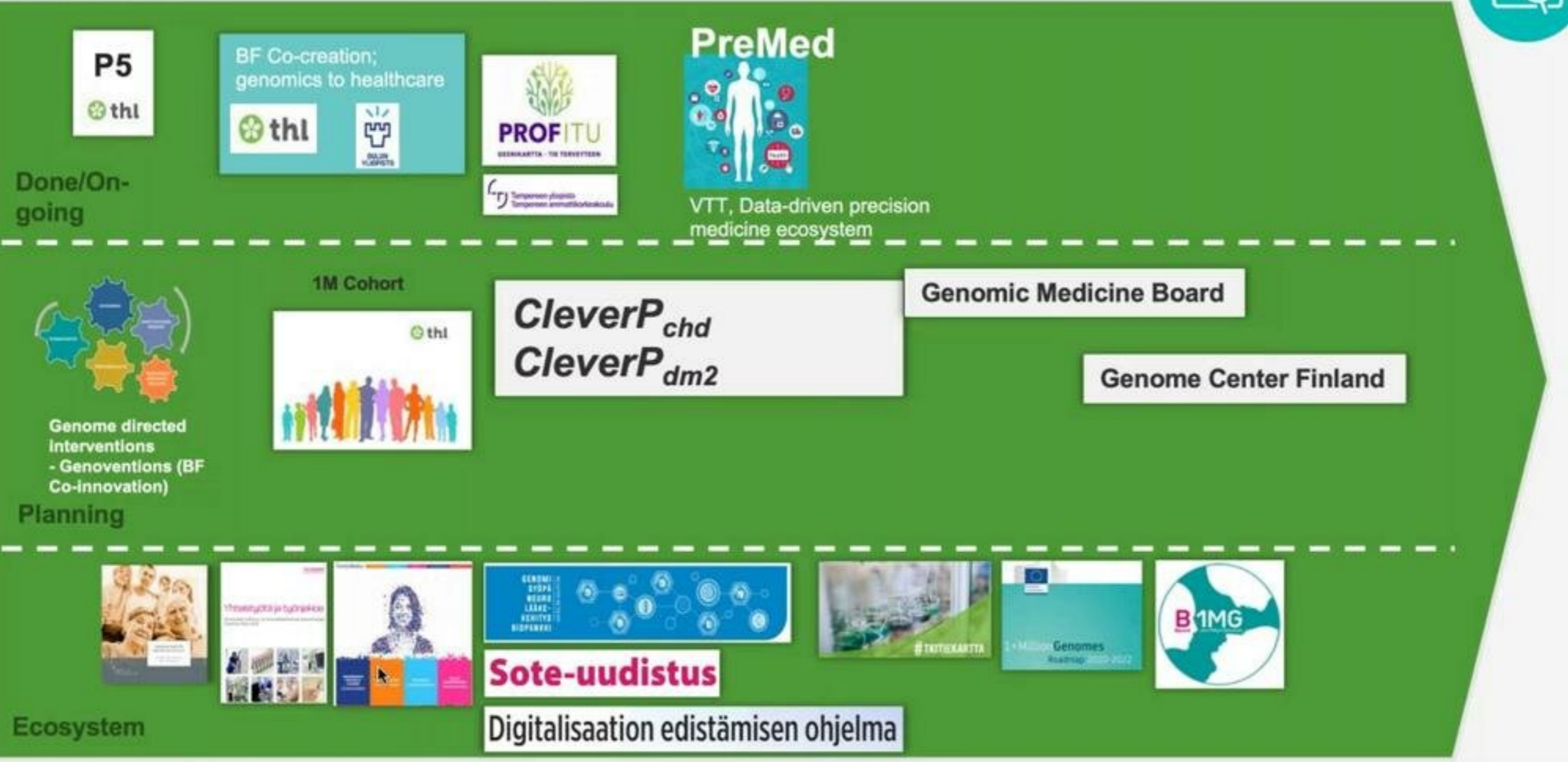


P6 – SUMMARY



1. **A large-scale study** returning information to 100.000 participants on their genetic risk for selected common chronic diseases.
 - 900.000 registry controls
2. A novel **operations model** for the healthcare professionals for using genetic information in their day-to-day clinical work
 - Predictive, preventive, personalized and participatory
 - Training and guidance to be provided
3. **Biobank portal**
4. **Novel opportunities** for research and public-private collaboration, incl. drug development and wellbeing services

Precision Public Health based on (Gen)omic data - P6 Genomics to Healthcare



P5 FINHEALTH PILOT



Polygenic risk score (PRS)

- Coronary artery disease
- T2 diabetes
- Venous thromboembolism

Single Clinical Variants (SCV)

The screenshot shows the MyP5 website interface. At the top, it displays the logo for 'TERVEYDEN JA HYVINVOINNIN LAITOS' and navigation links for 'Suomeksi', 'På Svenska', and 'in English'. Below the logo, there are navigation tabs: 'OmaP5', 'P5-tutkimus', 'Taudit', 'Monigeeninen riskisuunna', 'Elintavat', and 'Perinnöllisyys ja geenit'. The main content area features a login section titled 'Kirjautuminen' with fields for 'Käyttäjätunnus *' and 'Salasana *', a 'Kirjaudu sisään' button, and a 'Tulokset' link. To the right, there is a large image of an elderly man sitting on a rock, with a text overlay stating '~3500 participants'. Below the image, there is a 'Tervetulo' message and a list of features: 'Results portal' containing 'Results', 'Information', and 'Doctor's notes'. A small link 'Lisätunneille tietoa heidän' is visible at the bottom right.