Cost-Utility Analysis of Polygenic Risk Score in the Prevention of Type 2 Diabetes

Object-Oriented Individual-Level Model with Probabilistic Sensitivity Analysis (PSA)

Janne Martikainen¹ // Aku-Ville Lehtimäki¹ // Kari Jalkanen¹ // Piia Lavikainen¹ // Teemu Paajanen^{2,3} // Heidi Marjonen^{2,3} // Kati Kristiansson^{2,3} // Jaana Lindström² // Markus Perola^{2,3} ¹School of Pharmacy, University of Eastern Finland, Kuopio, Finland // ²Department of Public Health and Welfare, Finnish Institute for Health and Welfare, Helsinki, Finland // ³Research Program for Clinical and Molecular Metabolism, Faculty of Medicine, University of Helsinki, Finland

Background

- Type 2 diabetes (T2D) is a significant public health challenge.
- Obesity, unhealthy diet, and low physical activity are main risk factors of T2D, but genetic risk is also a factor in developing T2D.
- Recently, genomic-based polygenic risk scores (PRSs) have been suggested to improve the accuracy of risk prediction (1).
- The aim of this study was to assess the cost-utility of additional PRS testing (as a part of overall risk assessment) followed by a lifestyle intervention and an additional medical therapy when estimated 10year overall risk for T2D exceeded 20% among individuals screened as high-risk based on traditional risk factors only.

Methods

- For a cost-utility analysis, a microsimulation statetransition model with PSA was constructed.
- Model parameters were estimated based on the national FINRISK follow-up study (n=15,868) and other real-world datasets. For example, parametric survival regression models were used to assess the baseline risk of T2D with and without genetic risk information.
- The model was built utilizing object-oriented programming paradigm: an object was created to represent every individual virtually. In turn, each individual was assigned into a cohort object (Figure 1). The model was written in R (2).
- Cost-effectiveness acceptability curve (CEAC) and the expected value of perfect information (EVPI) were also estimated. Cohort-level PSA was used in evaluating the impact of simultaneous variation in model parameters on the model results.

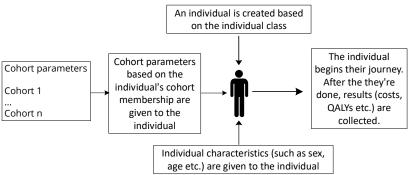


Figure 1. Overview of the object-oriented model structure

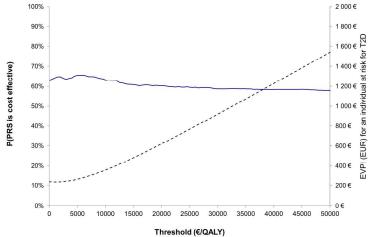


Figure 2. Cost-effectiveness acceptability curve (blue line; left y-axis) showing the probability that the PRS strategy is cost-effective compared to the usual practice, together with expected value of perfect information (dotted line; right y-axis) over a range of values for WTP

Results

- The use of targeted PRS strategy reclassified 12.4% of individuals as very high-risk individuals who would have been originally classified as high-risk.
- Over a lifetime horizon, the targeted PRS was the dominant strategy. The sensitivity analyses showed that PRS remained dominant in almost all simulations.
- However, there is still decision uncertainty since the probability of cost-effectiveness at a WTP of 0€/QALY was 63.0% (Figure 2). The corresponding EVPI estimate was 243€ per an individual.

Conclusions

 The results of the study (3) showed that PRS provides moderate additional value in risk screening leading to potential cost savings and better quality of life, comparing with the current T2D risk screening.

References:

- Mars N, Koskela JT, Ripatti P, Kiiskinen TTJ, Havulinna AS, Lindbohm JV, Ahola-Olli A, Kurki M, Karjalainen J, Palta P; FinnGen, Neale BM, Daly M, Salomaa V, Palotie A, Widén E, Ripatti S. Polygenic and clinical risk scores and their impact on age at onset and prediction of cardiometabolic diseases and common cancers. Nat Med. 2020;26(4):549-557.
- 2) R Core Team (2021). R: A language and environment for statistical computing. Boston: R
- Foundation for Statistical Computing. Available at: https://www.r-project.org/.

 3) Martikainen J, Lehtimäki AV, Jalkanen K, Lavikainen P, Paajanen T, Marjonen H, Kristiansson K, Lindström J, Perola M. Economic evaluation of using polygenic risk score to guide risk screening and interventions for the prevention of type 2 diabetes in individuals with high overall baseline risk. Front Genet. 2022 Sep 15;13:880799.





