CORRESPONDENCE



Genetics and Genomics

Response to: Genetic risk scores may compound rather than solve the issue of prostate cancer overdiagnosis (BJC-LT3342090)

© The Author(s), under exclusive licence to Springer Nature Limited 2022

British Journal of Cancer (2023) 128:487–488; https://doi.org/10.1038/s41416-022-02081-1

We thank the authors Horton et al. [1] for their comments on our recent paper about the potential application of a genetic risk score for prostate cancer detection in primary care settings [2]. We agree that the evidence base is not yet strong enough to support the use of genetic risk scores in clinic, although we would argue that Conti et al.'s GRS for prostate cancer [3] is reasonably, not weakly, predictive with an AUC of 0.703 when applied in our study.

The authors cite a study by Klein et al. as evidence that 'PSA is more predictive of prostate cancer metastasis or death than a genetic risk score' [4]. Klein et al. applied a risk score including 125 SNPs to two screening (asymptomatic) cohorts; the outcome was lifetime distant metastases or death from prostate cancer. In that study, the AUC of a single PSA test was 0.78, and the AUC of the GRS was 0.63. The AUC of GRS alone in our study was 0.703; the difference could be explained by the inclusion of a greater number of SNPs in the Conti score or the differences in study population. Crucially, our study included men with lower urinary tract symptoms and the outcome was prostate cancer within two years; this is different to the screening population.

The problem of false positives does deserve attention. The specificity of PSA is estimated at around 0.20; only one in five men who do not have prostate cancer will have a negative PSA test [5]. We hypothesise that combining GRS with PSA in clinic could reduce false positive referrals from primary care. Careful consideration of the appropriate threshold for an integrated risk model would be needed to ensure that more men are not 'swept along a pathway of ultimately unnecessary invasive investigations'. Currently, men who are referred get a pre-biopsy MRI, reducing the risk of unnecessary invasive investigations by 30–60%. This approach is less likely to find indolent prostate cancer. Changes in prostate biopsy from trans-rectal ultrasound to trans-perineal reduces risk of biopsy complications, and only 4% of men with indolent disease in the UK are over treated [6, 7].

The lack of ethnic diversity in our study remains a significant barrier for this and all research using the UK Biobank as a resource. Greater testing and evaluation of the use of GRS in primary care is needed to ensure that this approach results in health benefit, without increased risk of harm.

Harry D. Green¹, Samuel W. D. Merriel 60, Richard A. Oram³, Katherine S. Ruth⁴, Jessica Tyrrell⁴, Samuel E. Jones⁵, Chrissie Thirlwell^{6,7}, Michael N. Weedon³ and Sarah E. R. Bailey 60,

¹Exeter Centre of Excellence for Diabetes Research (EXCEED), University of Exeter Medical School, St Luke's Campus, University of Exeter, Heavitree Road, Devon, Exeter EX1 2LU, UK. ²DISCOVERY Group, University of Exeter Medical School, St Luke's Campus, University of Exeter, Heavitree Road, Devon, Exeter EX1 2LU, UK. ³Institute of Biomedical and Clinical Science, University of Exeter Medical School, St Luke's Campus, University of Exeter, Heavitree Road, Devon, Exeter EX1 2LU, UK. ⁴Genetics of Complex Traits, University of Exeter Medical School, University of Exeter, Exeter EX2 5DW, UK. ⁵Institute for Molecular Medicine (FIMM), University of Helsinki, Helsinki, Finland. ⁶University of Exeter Medical School, St Luke's Campus, University of Exeter, Heavitree Road, Devon, Exeter EX1 2LU, UK. ⁷UCL Cancer Institute, Huntley St, EX1 2LU London, UK. [∞]email: s.e.r.bailey@exeter.ac.uk

REFERENCES

- Rechtman L, Brenner S, Wright M, Ritsick M, Rahman F, Han M, et al. Genetic risk scores may compound rather than solve the issue of prostate cancer overdiagnosis. Br J Cancer. 2022;9:1692–701.
- Green HD, Merriel SWD, Oram RA, Ruth KS, Tyrrell J, Jones SE, et al. Applying a genetic risk score for prostate cancer to men with lower urinary tract symptoms in primary care to predict prostate cancer diagnosis: a cohort study in the UK Biobank. Br J Cancer. 2022;127:1534–9.
- Conti DV, Darst BF, Moss LC, Saunders EJ, Sheng X, Chou A, et al. Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nat Genet. 2021;53:413.
- Klein RJ. ARTICLE Prostate cancer polygenic risk score and prediction of lethal prostate cancer. Christel Häggström [Internet]. [cited 2022 Oct 19];18:17. Available from: https://doi.org/10.1038/s41698-022-00266-8.
- Merriel SWD, Pocock L, Gilbert E, Creavin S, Walter FM, Spencer A, et al. Systematic review and meta-analysis of the diagnostic accuracy of prostate-specific antigen (PSA) for the detection of prostate cancer in symptomatic patients. BMC Med. 2022;20:54.
- NICE guideline [NG131]. Overview. Prostate cancer: diagnosis and management. Guidance. NICE; 2019.
- National Prostate Cancer Audit. Results of the NPCA Prospective Audit in England and Wales for men diagnosed from 1 April 2019 to 31 March 2020 and the Impact of COVID-19 in England during 2020. London; 2022.

Received: 30 October 2022 Revised: 1 November 2022 Accepted: 17 November 2022

Published online: 19 December 2022

488

AUTHOR CONTRIBUTIONS

SB drafted the response. All authors reviewed before submission.

COMPETING INTERESTS

The authors declare no competing interests.

ADDITIONAL INFORMATION

Correspondence and requests for materials should be addressed to Sarah E. R. Bailey.

Reprints and permission information is available at http://www.nature.com/reprints

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.