Genetic Risk, Health-Associated Lifestyle, and Risk of Early-onset Total Cancer and Breast Cancer

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Abstract

Importance

Early-onset cancer (diagnosed under 50 years of age) is associated with aggressive disease characteristics and its rising incidence is a global concern. The association between healthy lifestyle and early-onset cancer and whether it varies by common genetic variants is unknown.

Objective

To examine the associations between genetic risk, lifestyle, and risk of early-onset cancers.

Design, Setting, and Participants

We analyzed a prospective cohort of 66,308 white British participants who were under age 50 and free of cancer at baseline in the UK Biobank.

Exposures

Sex-specific composite total cancer polygenic risk scores (PRSs), a breast cancer-specific PRS, and sexspecific health-associated lifestyle scores (HLSs, which summarize smoking status, body mass index [males only], physical activity, alcohol consumption, and diet).

Main Outcomes and Measures

Hazard ratios (HRs) and 95% confidence intervals (CIs) for early-onset total and breast cancer.

Results

A total of 1,247 incident invasive early-onset cancer cases (female: 820, male: 427, breast: 386) were documented. In multivariable-adjusted analyses with 2-year latency, higher genetic risk (highest vs. lowest tertile of PRS) was associated with significantly increased risks of early-onset total cancer in females (HR, 95% CI: 1.85, 1.50-2.29) and males (1.94, 1.45-2.59) as well as early-onset breast cancer in females (3.06, 2.20-4.25). An unfavorable lifestyle (highest vs. lowest category of HLS) was associated with higher risk of total cancer and breast cancer in females across genetic risk categories; the association with total cancer was stronger in the highest genetic risk category than the lowest: HRs in females and men were 1.85 (1.02, 3.36), 3.27 (0.78, 13.72) in the highest genetic risk category and 1.15 (0.44, 2.98), 1.16 (0.39, 3.40) in the lowest.

Conclusions and Relevance

Both genetic and lifestyle factors were independently associated with early-onset total and breast cancer risk. Compared to those with low genetic risk, individuals with a high genetic risk may benefit more from adopting a healthy lifestyle in preventing early-onset cancer.

Introduction

Cancers are caused by inherited variants and acquired mutations induced by environmental factors or result from unavoidable DNA replication errors. Modifiable lifestyle factors, such as smoking, physical activity, and diet, and their joint effect have been linked with the incidence of overall and major cancer types.^{1,2} In the United States, data from the Centers for Disease Control and Prevention and the National Cancer Institute suggested that over 40% of all incident cancers and cancer deaths were attributable to unhealthy lifestyle factors.³ Evidence from large population-based prospective studies suggest that a substantial number of cancer cases may be preventable through lifestyle modification.¹ On the other hand, genomewide association studies (GWAS) have identified thousands of susceptibility loci across major cancer types.⁴ Emerging studies have investigated the degree to which adherence to a healthy lifestyle may attenuate the risk of cancer across strata defined by common genetic variants, aggregated into cancer polygenic risk scores (PRS). These studies have focused on overall cancer⁵ and a few major cancer types.⁶

Early-onset cancer, defined as cancer diagnosed under age 50, represents a unique spectrum of malignancies^{15,16} and generally manifests with a more aggressive disease phenotype.^{15,16} The incidence of early-onset cancer has increased globally over the past decades, possibly supporting a basis in changing environmental hazards or interactions between hazardous environments and genetics.^{15,17-19} Mounting evidence has established close epidemiological and biological links between unhealthy lifestyle and early-onset cancer.²⁰ In addition, cancer PRS tend to be more strongly associated with early-onset compared to late-onset cancer.²¹ However, no previous study has evaluated the magnitude to which adopting a healthy lifestyle may attenuate the impact of common genetic variants on early-onset cancer risk, highlighting a significant knowledge gap.

To address this important unanswered question, we conducted a large prospective cohort study to investigate the association between genetic risk, health-associated lifestyle, and risk of early-onset total cancer in both sexes, as well as early-onset breast cancer in females (which accounts for almost half of the early-onset cancer cases in our female study population).

Study Population

The study was performed using data from the UK Biobank longitudinal cohort, the details of which have been described previously.²²⁻²⁴ In brief, the UK Biobank began between 2006 and 2010 when more than 500,000 participants aged 40 to 70 years from 22 assessment centers across England, Scotland, and Wales were enrolled. Information on demographics (age, sex, ethnicity), lifestyle and other health-related factors were collected via extensive baseline questionnaires, interviews, and physical measurements. Blood samples were collected at baseline and were used for genotyping. Informed consent was obtained from participants during the baseline assessment.

Ascertainment of Analytic Population

We restricted the study population to self-reported white British participants. We excluded participants diagnosed with any cancer before or at the cohort baseline, aged over 50 at the cohort baseline, or without genotype data. We further excluded individuals with genetic sex discordance, second (or higher)-degree related individuals (kinship coefficients >0.088), and heterozygosity or call rate outliers based on genotyping data. Individuals who had withdrawn consent to participate, or with missingness in smoking, body mass index (BMI), physical activity, alcohol intake, and diet were further removed, resulting in 66,308 eligible participants (females: 34,383, males: 31,925) for the final analyses.

Ascertainment of Genetic Risk

Genetic risk for early-onset total cancer was assessed using sex-specific composite PRSs. Briefly, we calculated composite PRSs as weighted sums of a spectrum of published cancer site-specific PRSs for European ancestry populations.²⁵⁻³⁴ These PRSs were selected based on their original training sample size, methods, test-set performance, possibility of overfit, and the availability of SNPs and weights. We used weights obtained from lasso regression by regressing early-onset total cancer on the PRSs of individual cancers (including lifestyle factors and a range of variables selected a priori [detailed in the Statistical Analysis section] as covariates, with a partial penalty on PRSs only).

We constructed sex-specific composite total cancer PRSs due to the difference in cancer spectra between females and males. Specifically in analyses focused on early-onset breast cancer among females, we constructed a breast cancer-specific PRS.²⁵ The lists of published cancer site-specific PRSs included in the lasso regression and those ultimately selected for the development of sex-specific composite total cancer PRSs are summarized in eTables 1-2 in Supplement 1. The complete lists of SNPs included in the cancer site-specific PRSs are presented in eTables S1-S23 in Supplement 2. The early-onset cancer spectrum in females and males, showcasing cancers with published site-specific PRSs qualified for inclusion in the lasso regression, are summarized in eTable 3 in the Supplement 1.

PRSs for individual cancer types were calculated as weighted sums of the effect allele dosage for selected SNPs for each individual assuming an additive model based on the formula $\beta_1 \times \text{SNP}_{i,1} + \beta_2 \times \text{SNP}_{i,2} + ... + \beta_j \times \text{SNP}_{i,j} + ... + \beta_n \times \text{SNP}_{i,n}$, where $\text{SNP}_{i,j}$ is the effect allele dosage for SNP *j* for individual *i*, β_j is perallele log odds ratio for SNP *j* for a specific cancer type, n is the total number of selected SNPs for the specific cancer type. The sex-specific composite total cancer PRSs were constructed based on the formula $h_1 \times \text{PRS}_{i,1} + h_2 \times \text{PRS}_{i,2} + \dots + h_k \times \text{PRS}_{i,k}$, where $\text{PRS}_{i,k}$ is the PRS for cancer type *k* for individual *i*, h_k is the weight for cancer type *k* obtained from lasso regression.

Ascertainment of Health-associated Lifestyle Scores

Sex-specific health-associated lifestyle scores (HLSs) were calculated based on a combination of baseline smoking status, BMI (males only), physical activity, alcohol consumption, and diet.⁵⁻⁷ BMI was not included as a component of female-specific HLS due to the widely recognized inverse association of BMI with the risk of early-onset/pre-menopausal breast cancer,³⁵ coupled with the fact that early-onset total cancer in our female study population was predominantly driven by early-onset breast cancer. Healthy lifestyles were defined as no current smoking, normal BMI (18.5 to 25 kg/m²), adequate physical activity (exercised for at least 75 minutes of vigorous activity per week or 150 minutes of moderate activity per week or an equivalent combination), no alcohol consumption, and healthy diet (consumed an increased amount of fruits, vegetables, whole grains, and a reduced amount of red meats and processed meats).^{5-7,36,37}

To streamline the delivery of the public health message in a simple way, participants received a score of 1 if they didn't meet the criterion for a specific lifestyle factor or 0 otherwise.⁵⁻⁷ We then added up the sum across the lifestyle components resulting in a final unweighted HLS ranging from 0-4 (females) or 0-5 (males), with higher scores indicating an unhealthier lifestyle. More details of the sex-specific HLSs can be found in eTable 4 in Supplement 1.

Ascertainment of Cancer and Death

Incident invasive cancer cases were ascertained via linkage to the National Health Service central registers and death registries in England, Wales, and Scotland. Cases were coded using the International Classification of Diseases (10th Revision). Early-onset cancer cases were defined as those diagnosed under the age of 50 years.¹⁷ Deaths were ascertained through linkage to death registries.

Statistical Analysis

The analyses of early-onset total cancer were conducted separately by sex, given the difference in cancer spectra between females and males. The analyses of early-onset breast cancer were restricted to females only. To minimize the potential impact of reverse causation, a latency period was introduced by excluding the first 2 years of follow-up.

Person-years of follow-up was calculated from the baseline until the date of diagnosis of any cancer (except non-melanoma skin cancer), death, loss to follow-up, or 50th birthday, whichever occurred earliest. Descriptive analyses were performed to assess population characteristics across categories of PRS and HLS. Multivariable Cox proportional hazards models with age as the timescale were used to estimate hazard ratios (HRs) and 95% confidence intervals (CIs) of early-onset total cancer and early-onset breast cancer across PRS and HLS groups (PRS: high, intermediate, vs. low, based on tertiles; HLS: unhealthy, intermediate, vs. healthy, based on the predefined cutoffs detailed in eTable 4 in Supplement 1) in females and males, as well as for their joint associations by using the collapsed categories. HRs and 95% CIs per 1-SD increase in PRS and per 1-unit increase in HLS were also estimated. Moreover, stratified analysis was performed to assess the association between HLS and early-onset total cancer and breast cancer risk within each PRS stratum. The proportional hazards assumption was tested using the likelihood ratio test to

compare models with and without product terms between exposures and log-transformed age. No violation of the assumption was detected.

Covariates were selected a priori. Multivariate analyses of early-onset total cancer were stratified by sex, and adjusted for the first 10 genetic principal components (continuous), genotyping batch (categorical), average total household income (less than 18,000, 18,000-30,999, 31,000-51,999, 52,000-100,000, greater than 100,000, prefer not to answer or do not know), education (college or university degree, some professional qualifications, secondary education, others, prefer not to answer or do not know), BMI (continuous, kg/m²; in analyses of females only), and family history of cancer (yes, no; family history of breast cancer was used in analyses of breast cancer). PRS and HLS were mutually adjusted.

Multivariable analyses of early-onset breast cancer were adjusted for the above-mentioned covariates, and were additionally adjusted for age at menarche (<12, 12-13, \geq 14 years, prefer not to answer or do not know), parity (nulliparous, parous, prefer not to answer or do not know), age at first live birth (<25, 25-29, \geq 30 years, prefer not to answer or do not know), oral contraceptive use (yes, no, prefer not to answer or do not know), menopausal status and hormone replacement therapy use (premenopausal, postmenopausal with hormone replacement therapy, postmenopausal without hormone replacement therapy, prefer not to answer or do not know).

Data analyses were performed using R software (version 4.3.1, https://www.r-project.org/), PLINK (version 2.0, https://www.cog-genomics.org/plink/2.0/),³⁸ and the Polygenic Score Catalog Calculator (version 2.0, https://github.com/PGScatalog/pgsc_calc).³⁹ Tests were two-sided, with *P* values <0.05 indicating statistical significance.

Results

Population Characteristics

A total of 1,247 incident invasive early-onset cancer cases (820 in females, 427 in males) were documented among 66,308 eligible participants during 329,509 person-years of follow-up (females: 168,897, males: 160,612), including 386 cases of incident early-onset breast cancer in females. The top three most common

incident invasive early-onset cancers documented were breast cancer, melanoma, and colorectal cancer in females, and melanoma, colorectal cancer, and prostate cancer in males. (eTable 3 in Supplement 1) The distribution of age, HLS and its components (smoking status, BMI, physical activity, alcohol consumption, and diet), total household income, and education did not vary appreciably across the PRSs of interest (female-specific total cancer PRS, male-specific total cancer PRS, and breast cancer PRS). Further, we observe no difference in age at menarche, parity, age at first live birth, oral contraceptive use, menopausal status, and hormone replacement therapy use across the PRSs. In contrast, individuals with higher PRSs (indicating higher genetic risk) in both sexes tended to have family history of cancer, and females with higher PRSs were more likely to have undergone mammography. (Tables 1, and eTable 5 in Supplement 1)

For both sexes, individuals with a higher HLS (indicating an unhealthier lifestyle) tended to have lower total household income and education levels and were more likely to have a family history of cancer. Among females, those with a higher HLS were less likely to undergo mammography or be premenopausal. Additionally, they were less likely to have experienced menarche between 12-13 years of age and to have their first birth before age 25 years. Moreover, they were more likely to use oral contraceptives. (eTables 6-7 in Supplement 1)

PRS and the Risk of Early-onset Total and Breast Cancer

In multivariable-adjusted analyses with 2-year latency, higher genetic risk (highest vs. lowest tertile of PRS) was associated with significantly increased risks of early-onset total cancer in females (female-specific composite PRS HR=1.85; 95% CI, 1.50-2.29), early onset total cancer in males (male-specific composite PRS HR=1.94; 95% CI, 1.45-2.59), and early-onset breast cancer in females (breast-cancer PRS HR=3.06; 95% CI, 2.20-4.25). The HR (95% CI) per 1-SD increase in PRS was 1.29 (1.19, 1.40) for early-onset total cancer in females, 1.36 (1.22, 1.53) for early-onset total cancer in males, and 1.58 (1.41, 1.78) for early-onset breast cancer in females. (Figure 1)

HLS and the Risk of Early-onset Total and Breast Cancer

Adopting an unhealthy lifestyle (highest vs. lowest category of HLS) showed suggestive associations with increased risks in females of early-onset total cancer (female-specific HLS HR=1.49; 95% CI, 0.99-2.25), and early-onset breast cancer (HR=1.78; 95% CI, 0.97-3.24), but not for early-onset total cancer in males (male-specific HLS HR=1.14; 95% CI, 0.67-1.95). The associations among females were statistically significant when examining effect estimates on the scale of per 1-unit increase for early-onset total cancer (HR=1.12; 95% CI, 1.01-1.24) and early-onset breast cancer (HR=1.17; 95% CI, 1.02-1.35), but not in males for early-onset total cancer (HR=1.01; 95% CI, 0.90-1.14). (Figure 2)

HLS with Early-onset Total and Breast Cancer Risk, Stratified by PRS Category

The HRs (95% CI) of early-onset total cancer in females and males, and early-onset breast cancer in females associated with adopting an unfavorable lifestyle (highest vs. lowest category of HLS) were 1.85 (1.02, 3.36), 3.27 (0.78, 13.72), and 1.67 (0.71, 3.90), respectively, in those with high genetic risk; 1.25 (0.60, 2.57), 1.11 (0.42, 2.89), and 1.66 (0.54, 5.11), respectively, in those with intermediate genetic risk; and 1.15 (0.44, 2.98), 1.16 (0.39, 3.40), and 2.10 (0.57, 7.75), respectively, in those with low genetic risk. On the scale of per 1-unit increase in HLS, the HRs (95% CI) of early-onset total cancer in females and males, and early-onset breast cancer in females were 1.20 (1.03, 1.40), 1.06 (0.87, 1.29), and 1.22 (1.00, 1.48), respectively, in those with high genetic risk; 1.00 (0.84, 1.19), 1.01 (0.81, 1.26), and 1.14 (0.88, 1.48), respectively, in those with intermediate genetic risk; and 1.14 (0.93, 1.40), 0.98 (0.77, 1.25), and 1.07 (0.76, 1.51), respectively, in those with low genetic risk. The *P* value for the 2-df interaction test on the log HR scale was 0.39 for early-onset total cancer in females, 0.77 for early-onset total cancer in males, and 0.11 for early-onset breast cancer in females. (Figure 3)

Joint Associations of PRS and HLS with Early-onset Total and Breast Cancer Risk

Compared to individuals with both low genetic risk (lowest tertile of PRS) and a favorable lifestyle (lowest category of HLS), those with both high genetic risk (highest tertile of PRS) and an unfavorable lifestyle (highest category of HLS) had increased risks of early-onset total cancer in females (HR=3.07; 95% CI, 1.64-5.78) and males (HR=2.18; 95% CI, 0.78-6.11), and significantly higher early-onset breast cancer risk in females (HR=4.11; 95% CI, 1.56-10.85). (Figure 4)

Discussion

Early-onset cancer is generally more aggressive compared to late-onset cancer with its rising incidence becoming a global concern. Deciphering the relationship between genetic risk, lifestyle modification, and risk of early-onset cancers may inform preventive strategies. In this large prospective cohort study among white British individuals, we present the first epidemiological evidence for this relationship for early-onset cancers.

Our results showed that genetic predisposition and lifestyle factors each demonstrated independent associations with the risk of early-onset total cancer and breast cancer. Stratified analyses by PRSs indicate that adopting a healthy lifestyle is likely beneficial for all individuals. Impressively, individuals with a high genetic risk may derive greater benefits from adopting a healthy lifestyle to prevent early-onset total cancer, compared to those with a low or intermediate genetic risk, although we were unable to detect statistically significant interaction between HLS and PRS mainly due to the limited number of early-onset cancer cases. These findings may inform preventive strategies for early-onset cancer in populations with varying genetic risk profiles.

To our knowledge, no previous study has investigated the extent to which adopting a healthy lifestyle could mitigate the impact of common genetic variants on the risk of early-onset cancers. A prior study of European ancestry population in the UK Biobank examined genetic risk and the benefits of adherence to a healthy lifestyle in relation to total cancer, irrespective of age at onset.⁵ Their findings demonstrated an additive interaction between genetic and lifestyle factors, indicating that individuals with a higher genetic risk may benefit more from lifestyle modification in relation to overall cancer risk in both sexes.⁵ Another study of invasive breast cancer among females of European ancestry in the UK Biobank reported that for premenopausal females, lifestyle intervention could have the greatest impact on those with a high genetic risk.⁶ Conversely, for postmenopausal females, lifestyle intervention might offer essentially similar benefits regardless of an individual's genetic predisposition.⁶ The associations of genetic risk and lifestyle with risk of other major cancers, including cancers of thyroid, lung, stomach, pancreas, colorectum, ovarian, kidney,

bladder, uterine, prostate, non-Hodgkin's lymphoma, lymphocytic leukemia, and melanoma, have been reported.⁸⁻¹⁴ However, no such evidence has been reported for early-onset cancers.

Strengths and Limitations

Our study has several strengths of note, including: 1) It provides the first epidemiological evidence addressing this important unanswered question in early-onset cancer prevention. 2) The prospective cohort study design, which minimized the potential for recall bias and selection bias. 3) The large sample size, with over 66,000 eligible participants involved in the final analytical population. 4) Ascertainment of cancer cases through linkage to national registries enhanced internal validity. 5) A wide spectrum of potential confounding factors was selected a priori and included as covariates in multivariable-adjusted analyses, which ensures relatively rigorous control for confounding. 6) Standardized protocols to assess heritable and lifestyle factors. 7) Novel approach for developing sex-specific composite total cancer PRSs to assess genetic risk of early-onset total cancer. 8) The concern of overfitting multicancer PRSs has been mitigated, given that 19 of the 23 site-specific cancer PRSs included did not incorporate UK Biobank data in their training samples. On average, the UK Biobank constitutes only 13% of the case data across these PRSs. 9) Applied 2-year latency to minimize the potential influence of reverse causation. 10) The analyses of early-onset total cancer were performed separately by sex to reflect the difference in cancer spectrum.

Our study also has several limitations: 1) The possibility for residual confounding cannot be completely ruled out due to the observational nature of study design. Making causal arguments should be approached with caution. 2) The aggregate analytic approach in analyses early-onset total cancer may potentially mask heterogeneity of effects across different cancer types, though it increases the power to detect the effects of interest. 3) The inability to explore heterogeneity across individual cancer types other than breast cancer further, due to power limitations. 4) The ethnic homogeneity of our analytic population (all white British) may limit generalizability of current findings to other groups. 5) Lifestyle factors were assessed based on a single measurement at baseline rather than through repeated measurements (which can better depict long-term trends), raising concern about the potential influence of random within-person variation. 6) We were not able to consider the age of 45 years⁴⁰ as the cutoff in analyses of early-onset breast cancer due to the

limited number of incident cases. We were unable to consider separate analysis by ER status because information on hormone-receptor subtype was unavailable in the UK Biobank.

Future large prospective investigations with repeated measurements and long-term follow-up are warranted to provide additional evidence on these associations in early-onset cancers in diverse populations and to examine potential heterogeneity across individual cancer types in the associations of interest.

Conclusions

Both genetic and lifestyle factors were independently associated with risks of early-onset total cancer and breast cancer. Compared to those with low genetic risk, individuals with a high genetic risk may benefit more from adopting a healthy lifestyle in preventing early-onset cancer.

Article Information

Author Contributions

YZ, YL and PK have full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. Concept and design: YZ, YL, and PK. Acquisition, analysis, or interpretation of data: All authors. Drafting of the manuscript: YZ and YL. Critical revision of the manuscript for important intellectual content: SL and PK. Statistical analysis: YZ and YL. Obtained funding: YZ, SL, and PK. Administrative, technical, or material support: SL and PK. Supervision: PK and YL.

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Conflict of Interest Disclosures

The authors declare no potential conflicts of interest.

Disclaimer

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Ethical Approval

The details of UK Biobank research ethics approval are elaborated at https://www.ukbiobank.ac.uk/learnmore-about-uk-biobank/about-us/ethics.

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Table 1. Characteristics of female and male participants at baseline according to female-specific and male-specific total cancer PRSs

	Females (N=34383)		Males (N=31925)			
Characteristics ^{ab}	Female-specific	Total Cancer PR	S	Male-specific Total Cancer PRS		
	Low (N=11444)	Intermediate (N=11476)	High (N=11463)	Low (N=10601)	Intermediate (N=10607)	High (N=10717)
Age, mean (SD)	45.0 (2.73)	45.0 (2.72)	45.0 (2.72)	44.9 (2.75)	45.0 (2.74)	44.9 (2.75)
Sex-specific HLS, n (%)				~ /	× ,	· · · · ·
Healthy	2566 (22.4%)	2515 (21.9%)	2507 (21.9%)	792 (7.5%)	740 (7.0%)	745 (7.0%)
Intermediate	8462 (73.9%)	8527 (74.3%)	8477 (74.0%)	6625 (62.5%)	6560 (61.8%)	6636 (61.9%)
Unhealthy	416 (3.6%)	434 (3.8%)	479 (4.2%)	3184 (30.0%)	3307 (31.2%)	3336 (31.1%)
Smoking, n (%)		~ /			× ,	· · · · ·
Current smoker	1204 (10.5%)	1216 (10.6%)	1225 (10.7%)	1551 (14.6%)	1547 (14.6%)	1568 (14.6%)
Others	10240 (89.5%)	10260 (89.4%)	10238 (89.3%)	9050 (85.4%)	9060 (85.4%)	9149 (85.4%)
BMI, n (%)		. ,	· · · ·			
$18.5 \le \text{to} < 25 \text{ kg/m}^2$	5447 (47.6%)	5477 (47.7%)	5538 (48.3%)	2912 (27.5%)	2914 (27.5%)	2964 (27.7%)
Others	5997 (52.4%)	5999 (52.3%)	5925 (51.7%)	7689 (72.5%)	7693 (72.5%)	7753 (72.3%)
Physical activity, n (%)						
Met the guidelines ^c	5901 (51.6%)	5877 (51.2%)	5838 (50.9%)	6224 (58.7%)	6127 (57.8%)	6183 (57.7%)
Didn't meet the guidelines	5543 (48.4%)	5599 (48.8%)	5625 (49.1%)	4377 (41.3%)	4480 (42.2%)	4534 (42.3%)
Alcohol intake, n (%)						
Never	637 (5.6%)	575 (5.0%)	567 (4.9%)	413 (3.9%)	423 (4.0%)	406 (3.8%)
Special occasions only	1274 (11.1%)	1311 (11.4%)	1222 (10.7%)	617 (5.8%)	567 (5.3%)	666 (6.2%)
One to three times a month	1745 (15.2%)	1646 (14.3%)	1790 (15.6%)	1184 (11.2%)	1217 (11.5%)	1226 (11.4%)
Once or twice a week	3432 (30.0%)	3403 (29.7%)	3533 (30.8%)	3318 (31.3%)	3236 (30.5%)	3286 (30.7%)
Three or four times a week	2818 (24.6%)	2818 (24.6%)	2737 (23.9%)	2977 (28.1%)	3015 (28.4%)	3057 (28.5%)
Daily or almost daily	1538 (13.4%)	1723 (15.0%)	1614 (14.1%)	2092 (19.7%)	2149 (20.3%)	2076 (19.4%)
Fruit and vegetable intake, n (%)						
<3 servings/day	3958 (34.6%)	4105 (35.8%)	4014 (35.0%)	4735 (44.7%)	4867 (45.9%)	4880 (45.5%)
\geq 3 to <5 servings/day	4966 (43.4%)	4906 (42.8%)	5025 (43.8%)	3886 (36.7%)	3867 (36.5%)	3849 (35.9%)
≥5 servings/day	2520 (22.0%)	2465 (21.5%)	2424 (21.1%)	1980 (18.7%)	1873 (17.7%)	1988 (18.6%)
Whole grain intake, n (%)						
<2 servings/day	7342 (64.2%)	7249 (63.2%)	7302 (63.7%)	5766 (54.4%)	5849 (55.1%)	5884 (54.9%)
≥ 2 to <5.5 servings/day	4062 (35.5%)	4187 (36.5%)	4120 (35.9%)	4525 (42.7%)	4454 (42.0%)	4542 (42.4%)
≥5.5 servings/day	40 (0.3%)	40 (0.3%)	41 (0.4%)	310 (2.9%)	304 (2.9%)	291 (2.7%)
Red and processed meat intake, n (%)						
<2 times/week	2330 (20.4%)	2362 (20.6%)	2274 (19.8%)	1033 (9.7%)	966 (9.1%)	986 (9.2%)
≥ 2 to <4 times/week	5478 (47.9%)	5400 (47.1%)	5416 (47.2%)	3847 (36.3%)	3807 (35.9%)	3848 (35.9%)
≥4 times/week	3636 (31.8%)	3714 (32.4%)	3773 (32.9%)	5721 (54.0%)	5834 (55.0%)	5883 (54.9%)
Average total household income, n (%)						
Less than 18,000	1199 (10.5%)	1181 (10.3%)	1208 (10.5%)	788 (7.4%)	833 (7.9%)	820 (7.7%)
18,000 to 30,999	1907 (16.7%)	1881 (16.4%)	1945 (17.0%)	1525 (14.4%)	1546 (14.6%)	1587 (14.8%)
31,000 to 51,999	3379 (29.5%)	3271 (28.5%)	3353 (29.3%)	3251 (30.7%)	3176 (29.9%)	3307 (30.9%)
52,000 to 100,000	3268 (28.6%)	3396 (29.6%)	3292 (28.7%)	3467 (32.7%)	3502 (33.0%)	3414 (31.9%)
Greater than 100,000	924 (8.1%)	937 (8.2%)	888 (7.7%)	1000 (9.4%)	981 (9.2%)	982 (9.2%)
Prefer not to answer or do not know	767 (6.7%)	810 (7.1%)	777 (6.8%)	570 (5.4%)	569 (5.4%)	607 (5.7%)
Family history of cancer, n (%)	0.405 (54.55)		0010 (70.000)			
No	8497 (74.2%)	8161 (71.1%)	8019 (70.0%)	7624 (71.9%)	7588 (71.5%)	7685 (71.7%)
Yes	2947 (25.8%)	3315 (28.9%)	3444 (30.0%)	2977 (28.1%)	3019 (28.5%)	3032 (28.3%)
Education, n (%)						

College or University degree	4626 (40.4%)	4741 (41.3%)	4644 (40.5%)	4276 (40.3%)	4127 (38.9%)	4218 (39.4%)
Some professional qualifications	297 (2.6%)	301 (2.6%)	309 (2.7%)	172 (1.6%)	197 (1.9%)	175 (1.6%)
Secondary education	6160 (53.8%)	6115 (53.3%)	6190 (54.0%)	5679 (53.6%)	5793 (54.6%)	5820 (54.3%)
Others	337 (2.9%)	294 (2.6%)	290 (2.5%)	437 (4.1%)	457 (4.3%)	466 (4.3%)
Prefer not to answer	24 (0.2%)	25 (0.2%)	30 (0.3%)	37 (0.3%)	33 (0.3%)	38 (0.4%)
Age at menarche, n (%)						
<12 years of age	1970 (17.2%)	1953 (17.0%)	1983 (17.3%)			
12 to 13 years of age	5054 (44.2%)	5088 (44.3%)	5015 (43.7%)			
\geq 14 years of age	4168 (36.4%)	4167 (36.3%)	4184 (36.5%)			
Prefer not to answer or do not know	252 (2.2%)	268 (2.3%)	281 (2.5%)			
Parity, n (%)						
Nulliparous	3027 (26.5%)	3057 (26.6%)	3112 (27.1%)			
Parous	8414 (73.5%)	8417 (73.3%)	8347 (72.8%)			
Prefer not to answer or do not know	3 (0.0%)	2 (0.0%)	4 (0.0%)			
Age at first live birth, n (%)						
<25 years of age	1981 (17.3%)	2054 (17.9%)	1955 (17.1%)			
25 to 29 years of age	2543 (22.2%)	2543 (22.2%)	2512 (21.9%)			
\geq 30 years of age	2090 (18.3%)	2056 (17.9%)	2074 (18.1%)			
Prefer not to answer or do not know	4830 (42.2%)	4823 (42.0%)	4922 (42.9%)			
Oral contraceptive use, n (%)						
Yes	10504 (91.8%)	10573 (92.1%)	10560 (92.1%)			
No	925 (8.1%)	894 (7.8%)	895 (7.8%)			
Prefer not to answer or do not know	15 (0.1%)	9 (0.1%)	8 (0.1%)			
Menopausal status and hormone						
replacement therapy use, n (%)						
Postmenopausal	258 (2.20/)	272 (2,40/)	254(2,20/)			
(with hormone replacement therapy)	238 (2.3%)	272 (2.4%)	234 (2.2%)			
Postmenopausal	509 (1 40/)	514(450/)	176(120(1))			
(without hormone replacement therapy)	308 (4.4%)	514 (4.5%)	470 (4.2%)			
Premenopausal	10663 (93.2%)	10673 (93.0%)	10715 (93.5%)			
Prefer not to answer or do not know	15 (0.1%)	17 (0.1%)	18 (0.2%)			
History of mammograms, n (%)						
Yes	2831 (24.7%)	2920 (25.4%)	3134 (27.3%)			
No	8575 (74.9%)	8523 (74.3%)	8287 (72.3%)			
Prefer not to answer or do not know	38 (0.3%)	33 (0.3%)	42 (0.4%)			

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; SD, standard deviation.

^a Characteristics of eligible female participants.

^b Mean (SD) for continuous variables and n (%) for categorical variables.

^c Met the guidelines of 150 minutes of moderate activity per week or 75 minutes of vigorous activity (or an equivalent combination).

A. Female-specific Total Cancer PRS and Early-onset Total Cancer in Females B. Male-specific Total Cancer PRS and Early-onset Total Cancer in Males C. Breast Cancer PRS and Early-onset Breast Cancer in Females **Female-specific** Male-specific 4.5-4.5-4.5-**Total Cancer PRS Total Cancer PRS Breast Cancer PRS** 4.0. 4.0 4.0-Low Low Low Intermediate Intermediate Intermediate 3.5 3.5 3.5 Multivariable-adjusted Hazard Ratios Multivariable-adjusted Hazard Ratios Multivariable-adjusted Hazard Ratios High High High Per 1-SD Per 1-SD Per 1-SD 2.5-2.5 2.0-20 1.5 0.5-0.5-0.5-0.0 0 0.0 P value <.50 <.001 <.001 .003 <.001 <.001 <.001 P value <.001 <.001 P value ---------

Figure legend

A: Multivariable-adjusted analysis of female-specific total cancer PRS and early-onset total cancer in females.

B: Multivariable-adjusted analysis of male-specific total cancer PRS and early-onset total cancer in males.

C: Multivariable-adjusted analysis of breast cancer PRS and early-onset breast cancer in females.

Figure 1. PRS and early-onset total and breast cancer risk

Multivariate analyses of early-onset total cancer were stratified by sex, and adjust for the first 10 genetic principal components for ancestry, genotyping batch, average total household income, education, BMI (females only), and family history of cancer (family history of breast cancer was used in analyses of breast cancer), plus adjustment of HLS. Multivariate analyses of early-onset breast cancer were adjusted for above-mentioned covariates, and were additionally adjusted for age at menarche, parity, age at first live birth, oral contraceptive use, menopausal status and hormone replacement therapy use, and history of mammograms. Reference group: individuals with low PRS.

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; SD, standard deviation; BMI, body mass index.



Figure 2: HLS and early-onset total and breast cancer risk

Figure legend

A: Multivariable-adjusted analysis of female-specific HLS and early-onset total cancer in females.

B: Multivariable-adjusted analysis of male-specific HLS and early-onset total cancer in males.

C: Multivariable-adjusted analysis of female-specific HLS and early-onset breast cancer in females.

Multivariate analyses of early-onset total cancer were stratified by sex, and adjust for the first 10 genetic principal components for ancestry, genotyping batch, average total household income, education, BMI (females only), and family history of cancer (family history of breast cancer was used in analyses of breast cancer), plus adjustment of PRS. Multivariate analyses of early-onset breast cancer were adjusted for above-mentioned covariates, and were additionally adjusted for age at menarche, parity, age at first live birth, oral contraceptive use, menopausal status and hormone replacement therapy use, and history of mammograms. Reference group: individuals with healthy HLS.

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; BMI, body mass index.



Figure 3: HLS and early-onset total and breast cancer risk, stratified by PRS category



Figure legend

A: Multivariable-adjusted analysis of female-specific HLS and early-onset total cancer in females, stratified by female-specific total cancer PRS.

B: Multivariable-adjusted analysis of male-specific HLS and early-onset total cancer in males, stratified by male-specific total cancer PRS.

C: Multivariable-adjusted analysis of female-specific HLS and early-onset breast cancer in females, stratified by breast cancer PRS.

Multivariate analyses of early-onset total cancer were stratified by sex, and adjust for the first 10 genetic principal components for ancestry, genotyping batch, average total household income, education, BMI (females only), and family history of cancer (family history of breast cancer was used in analyses of breast cancer). Multivariate analyses of early-onset breast cancer were adjusted for above-mentioned covariates, and were additionally adjusted for age at menarche, parity, age at first live birth, oral contraceptive use, menopausal status and hormone replacement therapy use, and history of mammograms. Reference group: individuals with healthy HLS within each PRS stratum.

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; BMI, body mass index.



Figure 4. Joint associations of PRS and HLS with early-onset total and breast cancer risk



Figure legend

A: Multivariable-adjusted joint association analysis of female-specific total cancer PRS and female-specific HLS with early-onset total cancer in females.

B: Multivariable-adjusted joint association analysis of male-specific total cancer PRS and male-specific HLS with early-onset total cancer in males.

C: Multivariable-adjusted joint association analysis of breast cancer PRS and female-specific HLS with earlyonset breast cancer in females.

Multivariate analyses of early-onset total cancer were stratified by sex, and adjust for the first 10 genetic principal components for ancestry, genotyping batch, average total household income, education, BMI (females only), and family history of cancer (family history of breast cancer was used in analyses of breast cancer). Multivariate analyses of early-onset breast cancer were adjusted for above-mentioned covariates, and were additionally adjusted for age at menarche, parity, age at first live birth, oral contraceptive use, menopausal status and hormone replacement therapy use, and history of mammograms. Reference group: individuals with both low PRS and healthy HLS.

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; BMI, body mass index.

Supplement 1

(Major Supplementary Tables)

Genetic Risk, Health-Associated Lifestyle, and Risk of Early-onset Total Cancer and Breast Cancer

Yin Zhang, Sara Lindström, Peter Kraft, Yuxi Liu

eTable 1

List of published cancer site-specific PRSs included in the lasso regression in females, and list of cancer sitespecific PRSs selected from the lasso regression for the development of the female-specific composite total cancer PRS

eTable 2

List of published cancer site-specific PRSs included in the lasso regression in males, and list of cancer sitespecific PRSs selected from the lasso regression for the development of the male-specific composite total cancer PRS

eTable 3

Summary of early-onset cancer spectrum in females and males, showcasing cancers with published site-specific PRSs qualified for inclusion in the lasso regression.

eTable 4

Components of female-specific HLS and male-specific HLS

eTable 5

Characteristics of female participants (N=34383) at baseline according to breast cancer PRS

eTable 6

Characteristics of female participants (N=34383) at baseline according to female-specific HLS

eTable 7

Characteristics of male participants (N=31925) at baseline according to male-specific HLS

eTable 1. List of published cancer site-specific PRSs included in the lasso regression in females, and list of cancer site-specific PRSs selected from the lasso regression for the development of the female-specific composite total cancer PRS

Cancer Types	Polygenic Score ID & Name (PGS catalog) included in the lasso regression ^a	Number of SNPs	Publication	PMID	Selected from the lasso regression
Breast	PGS000004 (PRS313_BC) ^b	313	Mavaddat N et al. Am J Hum Genet (2018)	30554720	Yes
Melanoma	PGS000790 (CC_Melanoma_IV)	24	Kachuri L et al. Nat Commun (2020)	33247094	No
Colorectum*	PGS002252 (PRS_CRC)	141	Archambault AN et al. J Natl Cancer Inst (2022) Huyghe JR et al. Nat Genet (2019)	35026030 30510241	No
Non-Hodgkin lymphoma	PGS000791 (CC_NHL_IV)	19	Kachuri L et al. Nat Commun (2020)	33247094	No
Ovary	PGS003394 (PRS Stepwise)	36	Dareng EO et al. Eur J Hum Genet (2022)	35027648	No
Brain [*]	PGS000624 (PRSWEB_PHECODE191.11_GWAS-Catalog- r2019-05-03-X191.11 PT UKB 20200608)	5	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Head and neck	PGS000792 (CC_Oral_IV)	14	Kachuri L et al. Nat Commun (2020)	33247094	No
Lung	PGS000740 (PRS128_LC)	128	Hung RJ et al. Cancer Res (2021)	33472890	Yes
Endometrium	PGS000786 (CC_Endo_IV)	9	Kachuri L et al. Nat Commun (2020)	33247094	No
Thyroid	PGS000797 (CC_Thyroid_IV)	12	Kachuri L et al. Nat Commun (2020)	33247094	No
Kidney	PGS000787 (CC_Kidney_IV)	19	Kachuri L et al. Nat Commun (2020)	33247094	No
Multiple myeloma*	PGS000653 (PRSWEB PHECODE204.4 GWAS-Catalog- r2019-05-03-X204.4 P 5e-08 UKB 20200608)	22	Fritsche LG et al. Am J Hum Genet (2020)	32991828	Yes
Cervix	PGS000784 (CC Cervix IV)	10	Kachuri L et al. Nat Commun (2020)	33247094	No

Pancreas	PGS000794 (CC_Pancreas_IV)	22	Kachuri L et al. Nat Commun (2020)	33247094	No
Stomach	PGS002299 (PRS3_gastric)	3	<u>Choi J et al. Int J</u> <u>Cancer (2020)</u> <u>Helgason et al. Nat</u> <u>Genet (2015)</u>	32588423 26098866	Yes
Lymphoid leukemia	PGS000788 (CC_LL_IV)	75	Kachuri L et al. Nat Commun (2020)	33247094	No
Esophagus	PGS000364 (PRSWEB PHECODE150 C15 LASSOSUM MGI 20200608)	2001	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Hodgkin's lymphoma	PGS000639 (PRSWEB_PHECODE201_GWAS-Catalog-r2019- 05-03-X201_PT_MGI_20200608)	20	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Trachea*	PGS000392 (PRSWEB_PHECODE165.1_GWAS-Catalog- r2019-05-03-X165.1_PT_UKB_20200608)	19	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No

Abbreviations: PRS, polygenic risk score; GRCh37, Genome Reference Consortium Human Build 37.

^a Published cancer site-specific PRSs included in the lasso regression were selected based on their training sample size, developing methods, test-set performance, possibility of overfit (i.e. training sample should not include UK Biobank), and the availability of SNPs and weights.

^b PGS000004 (PRS313_BC) was used specifically in analyses of breast cancer PRS and early-onset breast cancer among females.

^{*} UK Biobank was included in the training set.

eTable 2. List of published cancer site-specific PRSs included in the lasso regression in males, and list of cancer site-specific PRSs selected from the lasso regression for the development of the male-specific composite total cancer PRS

Cancer Types	Polygenic Score ID & Name (PGS catalog) included in the lasso regression ^a	Number of SNPs	Publication	PMID	Selected from the lasso regression
Breast	PGS000004 (PRS313_BC) ^b	313	Mavaddat N et al. Am J Hum Genet (2018)	30554720	No
Melanoma	PGS000790 (CC_Melanoma_IV)	24	Kachuri L et al. Nat Commun (2020)	33247094	Yes
Colorectum*	PGS002252 (PRS_CRC)	141	<u>Archambault AN et al. J</u> <u>Natl Cancer Inst (2022)</u> <u>Huyghe JR et al. Nat</u> <u>Genet (2019)</u>	35026030 30510241	No
Non-Hodgkin lymphoma	PGS000791 (CC_NHL_IV)	19	<u>Kachuri L et al. Nat</u> <u>Commun (2020)</u>	33247094	No
Prostate	PGS000662 (GRS.PCa.269)	269	Conti DV et al. Nat Genet (2021)	33398198	No
Brain*	PGS000624 (PRSWEB_PHECODE191.11_GWAS-Catalog- r2019-05-03-X191.11_PT_UKB_20200608)	5	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Head and neck	PGS000792 (CC Oral IV)	14	Kachuri L et al. Nat Commun (2020)	33247094	No
Lung	PGS000740 (PRS128_LC)	128	Hung RJ et al. Cancer Res (2021)	33472890	Yes
Testis	PGS000796 (CC_Testis_IV)	52	Kachuri L et al. Nat Commun (2020)	33247094	No
Thyroid	PGS000797 (CC_Thyroid_IV)	12	Kachuri L et al. Nat Commun (2020)	33247094	No
Kidney	PGS000787 (CC_Kidney_IV)	19	Kachuri L et al. Nat Commun (2020)	33247094	Yes
Multiple myeloma*	PGS000653 (PRSWEB_PHECODE204.4_GWAS-Catalog- r2019-05-03-X204.4_P_5e-08_UKB_20200608)	22	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Pancreas	PGS000794 (CC_Pancreas_IV)	22	Kachuri L et al. Nat Commun (2020)	33247094	Yes
Stomach	PGS002299 (PRS3 gastric)	3	Choi J et al. Int J Cancer (2020)	32588423 26098866	No

			Helgason et al. Nat Genet (2015)		
Lymphoid leukemia	PGS000788 (CC_LL_IV)	75	Kachuri L et al. Nat Commun (2020)	33247094	No
Bladder	PGS000782 (CC_Bladder_IV)	15	<u>Kachuri L et al. Nat</u> <u>Commun (2020)</u>	33247094	No
Esophagus	PGS000364 (PRSWEB_PHECODE150_C15_LASSOSUM_MGI_20200608)	2001	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Hodgkin's lymphoma	PGS000639 (PRSWEB_PHECODE201_GWAS-Catalog-r2019- 05-03-X201_PT_MGI_20200608)	20	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No
Larynx	PGS000362 (PRSWEB_PHECODE149.4_UKBB-SAIGE-HRC- X149.4_PT_MGI_20200608)	53	Fritsche LG et al. Am J Hum Genet (2020)	32991828	No

Abbreviations: PRS, polygenic risk score; GRCh37, Genome Reference Consortium Human Build 37.

^a Published cancer site-specific PRSs included in the lasso regression were selected based on their training sample size, developing methods, test-set performance, possibility of overfit (i.e. training sample should not include UK Biobank), and the availability of SNPs and weights.

^b Breast cancer PRS was included in the lasso regression in males because there was one case of early-onset breast cancer documented in males.

* UK Biobank was included in the training set.

eTable 3. Summary of early-onset cancer spectrum in females and males, showcasing cancers with published site-specific PRSs qualified for inclusion in the lasso regression

Cancer Types with Published Site-specific PRSs qualified for inclusion in the lasso regression	Number of Cases (jn females)	Number of Cases (in males)	Number of Cases (in the total study population)
Breast	386	1	387
Melanoma	60	36	96
Colorectum	23	28	51
Brain	11	19	30
Non-Hodgkin lymphoma	7	22	29
Prostate	0	28	28
Ovary	27	0	27
Testis	0	22	22
Kidney	7	9	16
Head and neck	7	9	16
Lung	8	7	15
Endometrium	12	0	12
Thyroid	9	3	12
Multiple myeloma	4	8	12
Lymphoid leukemia	4	5	9
Cervix	8	0	8
Pancreas	5	3	8
Stomach	2	5	7
Bladder	0	6	6
Esophagus	1	2	3
Hodgkin's lymphoma	2	1	3
Larynx	0	3	3
Trachea	1	0	1
Sum	584	217	801
Others	236	210	446
Total	820	427	1,247

eTable 4. Components of female-specific HLS and male-specific HLS

Components	Categories	Score assignments
Smoking status	Current smoker	Smoking score
	Others	1 for current smoker
		0 for others
BMI (males only) ^a	$18.5 \le \text{to} < 25 \text{ kg/m}^2$	BMI score
	Others	0 for $18.5 \le$ to $< 25 \text{ kg/m}^2$
		1 for others
Physical activity	Met the guidelines ^b	Physical activity score
	Didn't meet the guidelines ^b	0 for met the guidelines ^b
		1 for didn't meet the guidelines ^b
Alcohol consumption	Never	Alcohol intake score
	Others	0 for never
		1 for others
Diet	Fruit and vegetable intake	Fruit and vegetable intake
	<3 servings/day	1 for <3 servings/day
	\geq 3 to <5 servings/day	0.5 for \geq 3 to <5 servings/day
	≥5 servings/day	0 for \geq 5 servings/day
	*Amount per serving:	Whole grain intake
	Fresh fruit: 1 piece	1 for <2 servings/day
	Dried fruit: 5 pieces	0.5 for ≥ 2 to < 5.5 servings/day
	Cooked vegetables: 3 heaped tablespoons	0 for \geq 5.5 servings/day
	Salad/raw vegetables: 3 heaped tablespoons	
		Red and processed meat intake
	Whole grain intake	1 for ≥ 4 times/week
	<2 servings/day	0.5 for ≥ 2 to <4 times/week
	≥ 2 to <5.5 servings/day	0 for <2 times/week
	≥5.5 servings/day	
		Diet score
	*Amount per serving:	(Fruit and vegetable intake score +
	Cereal: 1 bowl	Whole grain intake score +
	Whole-meal bread: 1 slice	Red and processed meat intake score) / 3
	Whole-grain bread: 1 slice	1 for ≥ 0.5

Brown b	<i>read: 1 slice</i> 0 for <0.5
Red and ≥4 times ≥2 to <4 <2 times	processed meat intake /week times/week /week
	Total Score
<u>Female-specific HLS</u> = Smoking score + Physical activity score + Alcohol intake score + Diet score	<u>Male-specific HLS</u> = Smoking score + BMI score + Physical activity score + Alcohol intake score + Diet score
Ranging from 0 to 4, with higher scores indicating an unhealthin	er lifestyle. Ranging from 0 to 5, with higher scores indicating an unhealthier lifestyle.
Healthy:0 to 1Intermediate:2 to 3Unhealthy:4	Healthy:0 to 1Intermediate:2 to 3Unhealthy:4 to 5

Abbreviations: HLS, health-associated lifestyle score; BMI, body mass index.

^a BMI was not included as a component of female-specific HLS due to the widely recognized inverse association of BMI with the risk of early-onset/premenopausal breast cancer and the fact that early-onset total cancer in females was predominantly driven by early-onset breast cancer. ^b Met the guidelines of 150 minutes of moderate activity per week or 75 minutes of vigorous activity (or an equivalent combination).

	Breast Cancer PRS				
Characteristics ^{a,b}	Low	Intermediate	High		
	(N=11428)	(N=11481)	(N=11474)		
Age, mean (SD)	45.0 (2.72)	45.0 (2.73)	45.0 (2.72)		
Female-specific HLS, n (%)					
Healthy	2557 (22.4%)	2504 (21.8%)	2527 (22.0%)		
Intermediate	8468 (74.1%)	8524 (74.2%)	8474 (73.9%)		
Unhealthy	403 (3.5%)	453 (3.9%)	473 (4.1%)		
Smoking, n (%)					
Current smoker	1201 (10.5%)	1214 (10.6%)	1230 (10.7%)		
Others	10227 (89.5%)	10267 (89.4%)	10244 (89.3%)		
BMI, n (%)					
$18.5 \le \text{to} < 25 \text{ kg/m}^2$	5424 (47.5%)	5503 (47.9%)	5535 (48.2%)		
Others	6004 (52.5%)	5978 (52.1%)	5939 (51.8%)		
Physical activity, n (%)			× ,		
Met the guidelines ^c	5875 (51.4%)	5930 (51.7%)	5811 (50.6%)		
Didn't meet the guidelines	5553 (48.6%)	5551 (48.3%)	5663 (49.4%)		
Alcohol intake, n (%)			× ,		
Never	633 (5.5%)	591 (5.1%)	555 (4.8%)		
Special occasions only	1253 (11.0%)	1322 (11.5%)	1232 (10.7%)		
One to three times a month	1725 (15.1%)	1687 (14.7%)	1769 (15.4%)		
Once or twice a week	3418 (29.9%)	3431 (29.9%)	3519 (30.7%)		
Three or four times a week	2831 (24.8%)	2793 (24.3%)	2749 (24.0%)		
Daily or almost daily	1568 (13.7%)	1657 (14.4%)	1650 (14.4%)		
Fruit and vegetable intake, n (%)	× /				
<3 servings/day	3933 (34.4%)	4157 (36.2%)	3987 (34.7%)		
≥ 3 to <5 servings/day	4959 (43.4%)	4892 (42.6%)	5046 (44.0%)		
≥5 servings/day	2536 (22.2%)	2432 (21.2%)	2441 (21.3%)		
Whole grain intake, n (%)	× /				
<2 servings/day	7273 (63.6%)	7343 (64.0%)	7277 (63.4%)		
≥ 2 to <5.5 servings/day	4114 (36.0%)	4099 (35.7%)	4156 (36.2%)		
≥5.5 servings/day	41 (0.4%)	39 (0.3%)	41 (0.4%)		
Red and processed meat intake, n (%)	. ,				
<2 times/week	2344 (20.5%)	2310 (20.1%)	2312 (20.2%)		
≥ 2 to <4 times/week	5463 (47.8%)	5437 (47.4%)	5394 (47.0%)		
≥4 times/week	3621 (31.7%)	3734 (32.5%)	3768 (32.8%)		
Average total household income, n (%)			× ,		
Less than 18,000	1177 (10.3%)	1172 (10.2%)	1239 (10.8%)		
18,000 to 30,999	1888 (16.5%)	1922 (16.7%)	1923 (16.8%)		
31,000 to 51,999	3361 (29.4%)	3303 (28.8%)	3339 (29.1%)		
52,000 to 100,000	3311 (29.0%)	3361 (29.3%)	3284 (28.6%)		
Greater than 100,000	926 (8.1%)	923 (8.0%)	900 (7.8%)		
Prefer not to answer or do not know	765 (6.7%)	800 (7.0%)	789 (6.9%)		
Family history of breast cancer, n (%)					
No	10583 (92.6%)	10315 (89.8%)	10092 (88.0%)		
Yes	845 (7.4%)	1166 (10.2%)	1382 (12.0%)		
Education, n (%)			· /		
College or University degree	4649 (40.7%)	4702 (41.0%)	4660 (40.6%)		

eTable 5. Characteristics of female participants (N=34383) at baseline according to breast cancer PRS

Some professional qualifications	293 (2.6%)	305 (2.7%)	309 (2.7%)
Secondary education	6124 (53.6%)	6148 (53.5%)	6193 (54.0%)
Others	336 (2.9%)	302 (2.6%)	283 (2.5%)
Prefer not to answer	26 (0.2%)	24 (0.2%)	29 (0.3%)
Age at menarche, n (%)			
<12 years of age	1937 (16.9%)	1983 (17.3%)	1986 (17.3%)
12 to 13 years of age	5050 (44.2%)	5060 (44.1%)	5047 (44.0%)
\geq 14 years of age	4180 (36.6%)	4172 (36.3%)	4167 (36.3%)
Prefer not to answer or do not know	261 (2.3%)	266 (2.3%)	274 (2.4%)
Parity, n (%)			
Nulliparous	3059 (26.8%)	3026 (26.4%)	3111 (27.1%)
Parous	8366 (73.2%)	8453 (73.6%)	8359 (72.9%)
Prefer not to answer or do not know	3 (0.0%)	2 (0.0%)	4 (0.0%)
Age at first live birth, n (%)			
<25 years of age	2004 (17.5%)	2027 (17.7%)	1959 (17.1%)
25 to 29 years of age	2508 (21.9%)	2564 (22.3%)	2526 (22.0%)
\geq 30 years of age	2048 (17.9%)	2068 (18.0%)	2104 (18.3%)
Prefer not to answer or do not know	4868 (42.6%)	4822 (42.0%)	4885 (42.6%)
Oral contraceptive use, n (%)			
Yes	10473 (91.6%)	10595 (92.3%)	10569 (92.1%)
No	939 (8.2%)	878 (7.6%)	897 (7.8%)
Prefer not to answer or do not know	16 (0.1%)	8 (0.1%)	8 (0.1%)
Menopausal status and hormone			
replacement therapy use, n (%)			
Postmenopausal	266 (2.3%)	264(2.3%)	254(2.2%)
(with hormone replacement therapy)	200 (2.570)	204 (2.370)	254 (2.270)
Postmenopausal	504(4.4%)	512 (4 5%)	482 (4 2%)
(without hormone replacement therapy)	50+(+.+/0)	512 (4.570)	402 (4.270)
Premenopausal	10641 (93.1%)	10687 (93.1%)	10723 (93.5%)
Prefer not to answer or do not know	17 (0.1%)	18 (0.2%)	15 (0.1%)
History of mammograms, n (%)			
Yes	2799 (24.5%)	2954 (25.7%)	3132 (27.3%)
No	8592 (75.2%)	8490 (73.9%)	8303 (72.4%)
Prefer not to answer or do not know	37 (0.3%)	37 (0.3%)	39 (0.3%)

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; SD, standard deviation. ^a Characteristics of eligible female participants.

^b Mean (SD) for continuous variables and n (%) for categorical variables.

^c Met the guidelines of 150 minutes of moderate activity per week or 75 minutes of vigorous activity (or an equivalent combination).

	Female-specific HLS				
Characteristics ^{a,b}	Healthy	Intermediate	Unhealthy		
	(N=7588)	(N=25466)	(N=1329)		
Age, mean (SD)	45.0 (2.72)	45.0 (2.72)	45.0 (2.70)		
Female-specific Total Cancer PRS, n (%)					
Low	2566 (33.8%)	8462 (33.2%)	416 (31.3%)		
Intermediate	2515 (33.1%)	8527 (33.5%)	434 (32.7%)		
High	2507 (33.0%)	8477 (33.3%)	479 (36.0%)		
Smoking, n (%)		· · · ·			
Current smoker	36 (0.5%)	2280 (9.0%)	1329 (100%)		
Others	7552 (99.5%)	23186 (91.0%)	0 (0%)		
BMI. n (%)					
$18.5 \le \text{to} \le 25 \text{ kg/m}^2$	4204 (55.4%)	11712 (46.0%)	546 (41.1%)		
Others	3384 (44.6%)	13754 (54.0%)	783 (58.9%)		
Physical activity, n (%)		· · · ·			
Met the guidelines ^c	7279 (95.9%)	10337 (40.6%)	0(0%)		
Didn't meet the guidelines	309 (4.1%)	15129 (59.4%)	1329 (100%)		
Alcohol intake, n (%)		· · · ·			
Never	1132 (14.9%)	647 (2.5%)	0 (0%)		
Special occasions only	701 (9.2%)	2906 (11.4%)	200 (15.0%)		
One to three times a month	1008 (13.3%)	3974 (15.6%)	199 (15.0%)		
Once or twice a week	2211 (29.1%)	7822 (30.7%)	335 (25.2%)		
Three or four times a week	1772 (23.4%)	6307 (24.8%)	294 (22.1%)		
Daily or almost daily	764 (10.1%)	3810 (15.0%)	301 (22.6%)		
Fruit and vegetable intake, n (%)		× ,			
<3 servings/day	510 (6.7%)	10689 (42.0%)	878 (66.1%)		
≥ 3 to <5 servings/day	3252 (42.9%)	11240 (44.1%)	405 (30.5%)		
\geq 5 servings/day	3826 (50.4%)	3537 (13.9%)	46 (3.5%)		
Whole grain intake, n (%)					
<2 servings/day	3413 (45.0%)	17364 (68.2%)	1116 (84.0%)		
≥ 2 to <5.5 servings/day	4114 (54.2%)	8042 (31.6%)	213 (16.0%)		
≥5.5 servings/day	61 (0.8%)	60 (0.2%)	0 (0%)		
Red and processed meat intake, n (%)			· · ·		
<2 times/week	3294 (43.4%)	3572 (14.0%)	100 (7.5%)		
≥ 2 to <4 times/week	3723 (49.1%)	11961 (47.0%)	610 (45.9%)		
≥4 times/week	571 (7.5%)	9933 (39.0%)	619 (46.6%)		
Average total household income, n (%)					
Less than 18,000	771 (10.2%)	2576 (10.1%)	241 (18.1%)		
18,000 to 30,999	1183 (15.6%)	4274 (16.8%)	276 (20.8%)		
31,000 to 51,999	2131 (28.1%)	7489 (29.4%)	383 (28.8%)		
52,000 to 100,000	2317 (30.5%)	7362 (28.9%)	277 (20.8%)		
Greater than 100,000	625 (8.2%)	2072 (8.1%)	52 (3.9%)		
Prefer not to answer or do not know	561 (7.4%)	1693 (6.6%)	100 (7.5%)		
Family history of cancer, n (%)		. ,	. ,		
No	5536 (73.0%)	18226 (71.6%)	915 (68.8%)		
Yes	2052 (27.0%)	7240 (28.4%)	414 (31.2%)		
Education, n (%)					
College or University degree	3496 (46.1%)	10129 (39.8%)	386 (29.0%)		

eTable 6. Characteristics of female participants (N=34383) at baseline according to female-specific HLS

Some professional qualifications	208 (2.7%)	647 (2.5%)	52 (3.9%)
Secondary education	3674 (48.4%)	13993 (54.9%)	798 (60.0%)
Others	186 (2.5%)	645 (2.5%)	90 (6.8%)
Prefer not to answer	24 (0.3%)	52 (0.2%)	3 (0.2%)
Age at menarche, n (%)			
<12 years of age	1313 (17.3%)	4352 (17.1%)	241 (18.1%)
12 to 13 years of age	3376 (44.5%)	11233 (44.1%)	548 (41.2%)
\geq 14 years of age	2741 (36.1%)	9261 (36.4%)	517 (38.9%)
Prefer not to answer or do not know	158 (2.1%)	620 (2.4%)	23 (1.7%)
Parity, n (%)			
Nulliparous	2267 (29.9%)	6531 (25.6%)	398 (29.9%)
Parous	5317 (70.1%)	18933 (74.3%)	928 (69.8%)
Prefer not to answer or do not know	4 (0.1%)	2 (0.0%)	3 (0.2%)
Age at first live birth, n (%)			
<25 years of age	1202 (15.8%)	4450 (17.5%)	338 (25.4%)
25 to 29 years of age	1656 (21.8%)	5731 (22.5%)	211 (15.9%)
\geq 30 years of age	1340 (17.7%)	4756 (18.7%)	124 (9.3%)
Prefer not to answer or do not know	3390 (44.7%)	10529 (41.3%)	656 (49.4%)
Oral contraceptive use, n (%)			
Yes	6877 (90.6%)	23510 (92.3%)	1250 (94.1%)
No	702 (9.3%)	1933 (7.6%)	79 (5.9%)
Prefer not to answer or do not know	9 (0.1%)	23 (0.1%)	0 (0%)
Menopausal status and hormone			
replacement therapy use, n (%)			
Postmenopausal	168 (2.2%)	565 (2.2%)	51 (3.8%)
(with hormone replacement therapy)	100 (2.270)	505 (2.270)	51 (5.070)
Postmenopausal	339 (4 5%)	1069 (4.2%)	90 (6 8%)
(without hormone replacement therapy)	557 (4.570)	1009 (4.270)	90 (0.070)
Premenopausal	7070 (93.2%)	23796 (93.4%)	1185 (89.2%)
Prefer not to answer or do not know	11 (0.1%)	36 (0.1%)	3 (0.2%)
History of mammograms, n (%)			
Yes	1982 (26.1%)	6562 (25.8%)	341 (25.7%)
No	5580 (73.5%)	18821 (73.9%)	984 (74.0%)
Prefer not to answer or do not know	26 (0.3%)	83 (0.3%)	4 (0.3%)

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; SD, standard deviation. ^a Characteristics of eligible female participants.

^b Mean (SD) for continuous variables and n (%) for categorical variables.

^c Met the guidelines of 150 minutes of moderate activity per week or 75 minutes of vigorous activity (or an equivalent combination).

		Male-specific HLS		
Characteristics ^{a,b}	Healthy	Intermediate	Unhealthy	
	(N=2277)	(N=19821)	(N=9827)	
Age, mean (SD)	44.9 (2.76)	44.9 (2.74)	45.0 (2.74)	
Male-specific Total Cancer PRS, n (%)				
Low	792 (34.8%)	6625 (33.4%)	3184 (32.4%)	
Intermediate	740 (32.5%)	6560 (33.1%)	3307 (33.7%)	
High	745 (32.7%)	6636 (33.5%)	3336 (33.9%)	
Smoking, n (%)				
Current smoker	13 (0.6%)	1356 (6.8%)	3297 (33.6%)	
Others	2264 (99.4%)	18465 (93.2%)	6530 (66.4%)	
BMI, n (%)				
$18.5 \le \text{to} < 25 \text{ kg/m}^2$	2116 (92.9%)	6246 (31.5%)	428 (4.4%)	
Others	161 (7.1%)	13575 (68.5%)	9399 (95.6%)	
Physical activity, n (%)				
Met the guidelines ^c	2235 (98.2%)	14948 (75.4%)	1351 (13.7%)	
Didn't meet the guidelines	42 (1.8%)	4873 (24.6%)	8476 (86.3%)	
Alcohol intake, n (%)				
Never	407 (17.9%)	791 (4.0%)	44 (0.4%)	
Special occasions only	101 (4.4%)	1118 (5.6%)	631 (6.4%)	
One to three times a month	223 (9.8%)	2258 (11.4%)	1146 (11.7%)	
Once or twice a week	680 (29.9%)	6249 (31.5%)	2911 (29.6%)	
Three or four times a week	587 (25.8%)	5747 (29.0%)	2715 (27.6%)	
Daily or almost daily	279 (12.3%)	3658 (18.5%)	2380 (24.2%)	
Fruit and vegetable intake, n (%)		× ,	· · ·	
<3 servings/day	184 (8.1%)	7968 (40.2%)	6330 (64.4%)	
\geq 3 to <5 servings/day	959 (42.1%)	7676 (38.7%)	2967 (30.2%)	
≥5 servings/day	1134 (49.8%)	4177 (21.1%)	530 (5.4%)	
Whole grain intake, n (%)		× ,		
<2 servings/day	479 (21.0%)	10211 (51.5%)	6809 (69.3%)	
≥ 2 to <5.5 servings/day	1530 (67.2%)	9050 (45.7%)	2941 (29.9%)	
\geq 5.5 servings/day	268 (11.8%)	560 (2.8%)	77 (0.8%)	
Red and processed meat intake, n (%)				
<2 times/week	738 (32.4%)	1993 (10.1%)	254 (2.6%)	
≥ 2 to <4 times/week	1150 (50.5%)	7426 (37.5%)	2926 (29.8%)	
≥4 times/week	389 (17.1%)	10402 (52.5%)	6647 (67.6%)	
Average total household income, n (%)				
Less than 18,000	182 (8.0%)	1378 (7.0%)	881 (9.0%)	
18,000 to 30,999	344 (15.1%)	2949 (14.9%)	1365 (13.9%)	
31,000 to 51,999	658 (28.9%)	6140 (31.0%)	2936 (29.9%)	
52,000 to 100,000	720 (31.6%)	6408 (32.3%)	3255 (33.1%)	
Greater than 100,000	248 (10.9%)	1840 (9.3%)	875 (8.9%)	
Prefer not to answer or do not know	125 (5.5%)	1106 (5.6%)	515 (5.2%)	
Family history of cancer, n (%)		. /	. ,	
No	1659 (72.9%)	14243 (71.9%)	6995 (71.2%)	
Yes	618 (27.1%)	5578 (28.1%)	2832 (28.8%)	
Education, n (%)			· /	
College or University degree	1216 (53.4%)	7960 (40.2%)	3445 (35.1%)	

eTable 7. Characteristics of male participants (N=31925) at baseline according to male-specific HLS

Some professional qualifications	28 (1.2%)	332 (1.7%)	184 (1.9%)
Secondary education	970 (42.6%)	10666 (53.8%)	5656 (57.6%)
Others	58 (2.5%)	779 (3.9%)	523 (5.3%)
Prefer not to answer	5 (0.2%)	84 (0.4%)	19 (0.2%)

Abbreviations: PRS, polygenic risk score; HLS, health-associated lifestyle score; SD, standard deviation. ^a Characteristics of eligible male participants.

^b Mean (SD) for continuous variables and n (%) for categorical variables.

^c Met the guidelines of 150 minutes of moderate activity per week or 75 minutes of vigorous activity (or an equivalent combination).