ORIGINAL RESEARCH

Association between baseline serum glucose, triglycerides and total cholesterol, and prostate cancer risk categories

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Introduction

An increasing number of studies suggest a role for lipid and glucose metabolism in prostate cancer (PCa) development. Findings from a recent meta-analysis reported a

Abstract

Lifestyle-related risk factors such as hyperglycemia and dyslipidemia have been associated with several cancers. However, studies exploring their link with prostate cancer (PCa) clinicopathological characteristics are sparse and inconclusive. Here, we investigated the associations between serum metabolic markers and PCa clinicopathological characteristics. The study comprised 14,294 men from the Swedish Apolipoprotein MOrtality RISk (AMORIS) cohort who were diagnosed with PCa between 1996 and 2011. Univariate and multivariable logistic regression were used to investigate the relation between glucose, triglycerides and total cholesterol and PCa risk categories, PSA, Gleason score, and T-stage. Mean age at time of PCa diagnosis was 69 years. Men with glucose levels >6.9 mmol/L tend to have PSA<4 μ g/L, while those with glucose levels of 5.6–6.9 mmol/L had a greater odds of PSA>20 μ g/L compared to PSA 4.0–9.9 μ g/L. Hypertriglyceridemia was also positively associated with PSA>20 µg/L. Hyperglycemic men had a greater odds of intermediate- and high-grade PCa and advanced stage or metastatic PCa. Similarly, hypertriglyceridemia was positively associated with high-grade PCa. There was also a trend toward an increased odds of intermediate risk localized PCa and advanced stage PCa among men with hypertriglyceridemia. Total cholesterol did not have any statistically significant association with any of the outcomes studied. Our findings suggest that high serum levels of glucose and triglycerides may influence PCa aggressiveness and severity. Further investigation on the role of markers of glucose and lipid metabolism in influencing PCa aggressiveness and severity is needed as this may help define important targets for intervention.

> strong positive association between obesity and risk of advanced prostate cancer [1], indicating that lifestyle-related risk factors influence PCa aggressiveness and progression. Nonetheless, epidemiological evidence on the association between other lifestyle-related risk factors, including

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dyslipidemia and diabetes, and PCa development and progression remains sparse and inconclusive [2–5]. Moreover, most of these studies are limited by few cases, short follow-up time, and lack of power to detect the true associations between the exposures and the outcomes. Biologically, both dyslipidemia and hyperglycemia have been implicated with prostate carcinogenesis. Evidence from experimental studies using in vivo and in vitro models demonstrated that they may induce prostate carcinogenesis by modulating signaling pathways, which promote carcinogenic processes such as cell growth and proliferation, inflammation, oxidative stress, and cell migration [6–15].

Based on data in the AMORIS cohort, a large Swedish database with information on over 800,000 men and women, we have previously identified that abnormal serum lipid and glucose profiles may be involved in risk of developing PCa. Our findings showed a positive association between hypertriglyceridemia and PCa risk in hyper-glycemic men [16]. Conversely, we found that high-density lipoprotein and Apolipoprotein A-1 were inversely associated with PCa risk [17]. Here, we further investigated this by evaluating the association between serum total cholesterol, triglycerides and glucose and PCa risk categories and clinicopathological characteristics (i.e., PSA, Gleason score, and TNM staging) in the updated AMORIS cohort.

Materials and Methods

Study design and population

Detailed description of the AMORIS cohort can be found elsewhere [16, 18]. Briefly, this database comprises 812,073 Swedish men and women with blood samples sent for laboratory evaluation to the Central Automation Laboratory (CALAB) in Stockholm, Sweden, during the period 1985-1996 [19-21]. Individuals recruited were primarily from the greater Stockholm area, who were either healthy and having laboratory testing as part of a general checkup or outpatients referred for laboratory testing. None of the participants were inpatients at the time of sampling. In the AMORIS cohort, the CALAB database was linked to several Swedish national registries such as the Swedish National Cancer Register, the Hospital Discharge Register, the Cause of Death Register, the consecutive Swedish Censuses during 1970-1990, and the National Register of Emigration using the Swedish 10-digit personal identity number.

For this study, we specifically focused on the linkage between the AMORIS database and the National Prostate Cancer Register (NPCR), which has been nationwide since 1998 [22]. NPCR was developed to provide data for quality assurance and includes 98% of all newly diagnosed PCa cases registered in the Swedish National Cancer Register [22] to which reporting is mandated. From the NPCR, we extracted information on date of diagnosis, age at diagnosis, TNM stage [19], Gleason score, serum concentration of PSA at time of diagnosis, and primary treatment given or planned up to 6 months after date of diagnosis. Information on educational level was retrieved from the Population and Housing Census for 1970–1990. Using information from the National Patient Register, we calculated the Charlson comorbidity index which includes 19 diseases, with each disease category assigned a weight. The sum of an individual's weights was used to create a score, resulting in four comorbidity levels ranging from no comorbidity to severe comorbidity $(0, 1, 2, \text{ and } \geq 3)$ [23].

From AMORIS, we selected all men aged 20 and older diagnosed with PCa (and information on their prostate tumor characteristics from 1996 until 31 December 2011), who had baseline levels of serum glucose, total cholesterol, and triglycerides taken from the same health visit [19]. A total of 14,294 men were included in the final analysis. Figure S1 provides an overview of the cohort selection process.

Total cholesterol and triglycerides were measured enzymatically as previously described [24]. Glucose was also measured enzymatically with a glucose oxidase/peroxidase method. All methods were fully automated with automatic calibration and performed at one accredited laboratory [20].

The primary endpoints for our study are PCa risk categories. These risk categories were defined in accordance with an adapted version of National Comprehensive Cancer Network as low risk (T1-2, Gleason score 2-6 and PSA <10 ng/mL), intermediate risk (T1-2, Gleason score 7 and/or PSA 10 to <20 ng/mL), high risk (T3 and/or Gleason score 8-10 and/or PSA 20 to <50 ng/mL), and regionally metastatic tumors (T4 and/or N1 and/or PSA 50-<100 ng/mL in the absence of distant metastases (M0 or MX)) and distant metastatic tumors (M1 and/or PSA >100 ng/mL) [22]. We also included the clinicopathological characteristics (PSA levels, Gleason score, T stage) as outcomes. PSA was categorized as <4 µg/ L, 4-9.9 µg/ L, 10–20 μ g/L and >20 μ g/L. Gleason score was defined as low-grade tumor (Gleason score ≤6 or WHO grade 1), intermediate grade tumor (Gleason score 7 or WHO grade 2) and high-grade tumor (Gleason score ≥ 8 or WHO grade 3), while T stage was defined as localized (T1-T2) and advanced (T3-T4).

Statistical analysis

Odds ratios and 95% confidence intervals for the association between serum glucose, total cholesterol and triglycerides and PCa clinicopathological characteristics were calculated using logistic regression. In multivariable analyses, models were adjusted for age at diagnosis, educational level (low, intermediate, high), CCI (0, 1, 2, \geq 3), and fasting status (fasting, nonfasting, missing). For each man, we also calculated the interval time between the time from blood analyses and date of PCa diagnosis, which was also included as a covariate in the model. All models were also mutually adjusted for serum glucose, total cholesterol, and triglycerides. Due to the high percentage of cases with missing information on body mass index (BMI) (78.5%), BMI was not included as a covariate in our multivariate model. These missing value indicators were included in the statistical models. We did conduct a sensitivity analysis in the subgroup with information on BMI (n = 3073), whereby we adjusted for BMI. Due to the small sample size, statistical significance disappeared, but the directions of the associations virtually remained the same. We also conducted stepwise regression analyses among those with information on BMI to assess the effect of excluding BMI on the multivariate model. The results of our stepwise regression analyses showed that adjusting for BMI had little or no effect on the multivariate model, thus suggesting that BMI may not be an important confounder in this study population. Hence, all results shown are focused on the cohort of 14,294 men. Missing values were assigned to separate categories for education (1.5%) and fasting status (11.0%).

Serum glucose, total cholesterol, and triglycerides levels were analyzed using clinical cut-offs in accordance with the American Diabetes Association and National Cholesterol Education Programme (NCEP) guidelines[25–27]. Serum glucose levels were categorized as <5.6 mmol/L, 5.6–6.9 mmol/L, and >6.9 mmol/L, while serum total cholesterol was classified as <5.18 mmol/L, 5.18 mmol/L–6.19 mmol/L, and >6.19 mmol/L and serum triglycerides as <1.7 mmol/L, 1.7–2.24 mmol/L, and >2.24 mmol/L. The lowest clinical cut-offs were selected as the reference category.

Finally, we conducted a sensitivity analysis whereby we excluded men with measurements taken 2 years or less prior to the date of PCa diagnosis to assess possible reverse causation. We also performed a sensitivity analysis whereby we excluded men with nonfasting glucose measurements.

Data management and statistical analyses were conducted with Statistical Analysis Software (SAS) release 9.4 (SAS Institute, Cary, NC).

Results

Table 1 illustrates the baseline characteristics of the study population. Mean age at PCa diagnosis was 69 ± 8.0 years.

The median levels of glucose, total cholesterol, and triglycerides at baseline were 4.9 mmol/L, 5.8 mmol/L, and 1.2 mmol/L, respectively. Most men had normal glucose (<5.60 mmol/L) and triglycerides (<1.7 mmol/L) levels (82.5% and 69.1%, respectively), while at least 75% men had borderline high or high total cholesterol levels with mean time between measurement and PCa diagnosis of 16.7 \pm 4.8 years.

Men with glucose in the prediabetic range (5.6-6.9 mmol/L) had a greater odds of high risk and metastatic cancer, as compared to normoglycemic men (OR: 1.13; 95% CI: 0.98-1.30 and OR: 1.21; 95% CI: 1.04-1.42 for high risk and metastatic PCa, respectively) (Table 2). Triglycerides were positively associated with intermediate risk localized PCa for the highest versus the lowest subgroup (OR: 1.17; 95% CI: 1.03-1.34). Positive but nonstatistically significant associations between hypertriglyceridemia and high risk and metastatic PCa were also observed. When considering total cholesterol, there was no evidence to suggest any differences in any of the outcomes studies based on serum total cholesterol levels

When studying PSA as an outcome, we observed that a greater proportion of men with glucose levels >6.90 mmol/L had PSA <4 μ g/L (OR: 1.43; 95% CI: 1.01–1.45), compared to PSA levels of 4.0–9.9 μ g/L. There was also a positive association between glucose levels of 5.6–6.9 mmol/L and PSA levels >20 μ g/L compared to PSA levels of 4.0–9.9 μ g/L (OR: 1.28; 95% CI: 1.13–1.45) (Table 3). Moreover, we found a positive association between triglycerides ≥2.25 mmol/L and PSA levels >20 μ g/L (OR: 1.17; 95% CI: 1.03–1.34), as compared to PSA levels of 4.0–9.9 μ g/L.

For Gleason score, we found a positive association between glucose levels of 5.6-6.9 mmol/L and risk of intermediate-grade (OR: 1.18; 95% CI: 1.05-1.33) and high-grade PCa (OR: 1.24; 95% CI: 1.07-1.44) (Table 3). Compared to men with glucose <5.6 mmol/L, there was also an increased odds of high-risk PCa among men with glucose levels in the diabetic range (>6.9 mmol/L) (OR: 1.37; 95%: 1.02-1.87). Hypertriglyceridemia was positively associated with high-grade PCa (OR: 1.17; 95% CI: 1.01-1.37), but the odds of intermediate grade PCa did not appear to be different by triglycerides levels. Although neither glucose nor triglycerides had a statistically significant association with stage of PCa, there was an observed trend toward increased odds of T3-T4 PCa with increasing glucose and triglycerides levels (P for trend: 0.022 and 0.046, respectively).

The exclusion of 21 men with measurements taken less than 2 years prior to PCa diagnosis did not result in changes in the findings (Table S2 and S3). When we restricted our analyses to men with fasting glucose

Table 1. Baseline characteristics of study population.

Characteristics	Total population ($N = 14294$)
Age at diagnosis (years)	
Mean (SD)	69 (8.0)
N (%)	
≤49	95 (0.7)
50–59	1971 (13.8)
60–69	6321 (44.2)
≥70	5916 (41.4)
Education	
High	4427 (31.0)
Intermediate	5912 (41.2)
Low	3740 (26.2)
Missing	215 (1.5)
Charlson comorbidity index	
0	10651 (74.5)
1	1840 (12.9)
2	1104 (7.7)
≥3	699 (4.9)
Glucose (mmol/L)	
Median (IQR)	4.9 (4.6–5.4)
N (%)	/
<5.60	11796 (82.5)
5.6–6.9	2062 (14.4)
>6.9	436 (3.1)
Total cholesterol (mmol/L)	
Median (IQR)	5.8 (5.2–6.5)
N(%)	
<5.18	3552 (24.9)
5.18–6.19	5394 (37.7)
>6.19	5348 (37.4)
Triglycerides (mmol/L)	33.6(37.1)
Median (IQR)	1.2 (0.9–1.8)
N (%)	
<1.70	9878 (69.1)
1.70–2.24	2199 (15.4)
≥2.25	2217 (15.5)
Fasting status	22.7 (19.9)
N (%)	
Fasting	8577 (60.0)
Nonfasting	4143 (29.0)
Missing/Unknown	1574 (11.0)
5	ment and prostate cancer diagnosis
Mean (SD)	16.7 (4.8)
PSA (μ g/L)	10.7 (4.0)
Median (IQR)	9.4 (5.8–20.0)
N(%)	5.4 (5.6-20.0)
>4	1164 (8.1)
4.0–9.9	6126 (42.9)
4.0–9.9 10.0–20.0	3219 (22.5)
>20.0	3351 (23.4)
Missing Glasson grada	434 (3.0)
Gleason grade	
N (%)	(0.42)(42,5)
Low	6942 (48.6)
Intermediate	4789 (33.5)
High	2320 (16.2)
Missing	243 (1.7)

(Continued)

Table 1	(Continued)
Table L	

Characteristics	Total population ($N = 14294$)
Clinical stage	
N (%)	
Τ1	6797 (47.6)
T2	4290 (30.0)
Т3	2496 (17.5)
Τ4	319 (3.2)
Missing	392 (2.7)
NO	1502 (10.5)
N1	292 (1.0)
Missing/NX	12500 (87.5)
MO	4665 (32.6)
M1	867 (6.1)
Missing/MX	8762 (61.3)
Risk categories	
N (%)	
Localized	
Low risk	4472 (31.3)
Intermediate risk	4032 (28.2)
High risk	3258 (22.8)
Regional/distant metastatic	2187 (15.3)
Missing	345 (2.4)

measurements, the direction of the associations remained virtually unchanged (Table S4).

Discussion

In this study, we found evidence suggesting an independent positive association between serum glucose levels and higher grade (Gleason score 7 or ≥8) and advanced PCa (T3–T4, metastatic PCa). Similarly, men with hypertriglyceridemia (≥2.25 mmol/L) had increased odds of higher grade or more advanced PCa. When PSA was taken into account, we observed that men with glucose levels in the diabetic range (>6.9 mmol/L) had greater odds of having PSA levels <4 µg/L, while those with glucose levels in the prediabetic range (5.6–6.9 mmol/L) had greater odds of having PSA >20 µg/L. Men with hypertriglyceridemia (≥2.25 mmol/L) also had increased odds of PSA >20 µg/L. There was, however, no evidence to suggest that the odds of any of the outcomes studied varied by total cholesterol levels.

Glucose

Hyperglycemia has been positively associated with cancers such as pancreatic, breast, and colorectal cancer [28,29]. However, its link with prostate carcinogenesis is conflicting. Consistent with our findings, four other studies found evidence of higher risk of more aggressive or advanced PCa among men with abnormal glucose levels with the association being nonsignificant in two of the studies [3,

	Glucose (mmol/L) ¹	mol/L) ¹		Dfor	Total choles	Total cholesterol (mmol/L)		D for	Triglycerides (mmol/ L)	s (mmol/ L)		Dfor
	<5.60	5.60-6.90	>6.90	trend	<5.18	5.18-6.19	>6.19	trend	<1.70	1.70–2.24	≥2.25	trend
Low risk (Reference level for outcome studied) N = 4472 3794 558	level for outc 3794	ome studied) 558	120		1189	1689	1594		3152	690	630	
N = 4032 Age-adjusted OR	3359 1.00 (Ref)	566 1.08 (0.95–1.23)	107 0.94 (0.72–1.22)	0.601	1029 1.00 (Ref)	1527 0.97 (0.87–1.08)	1476 0.96 (0.86–1.07)	0.469	2784 1.00 (Ref)	603 0.99 (0.87–1.11)	645 1.17 (1.03–1.32)	0.030
(95% CI) Multivariable adjusted OR (95% CI)	1.00 (Ref)	1.00 (Ref) 1.07 (0.94–1.21) 0.90 (0.69–1.19)		0.864	1.00 (Ref)	0.96 (0.86–1.07)	0.93 (0.83–1.05)	0.228	1.00 (Ref)	0.99 (0.88–1.12)	0.99 (0.88–1.12) 1.17 (1.03–1.34)	0.037
High risk N = 3258 Age-adjusted OR	2630 1.00 (Ref)	523 105 1.23 (1.07–1.41) 1.09 (0.82–1.44)		0.016	752 1.00 (Ref)	1216 1.00 (0.88–1.13)	1290 1.06 (0.94–1.20)	0.287	2216 1.00 (Ref)	529 1.09 (0.96–1.25)	513 1.20 (1.05–1.37)	0.006
עור אלפט) Multivariable adjusted OR	1.00 (Ref)	1.00 (Ref) 1.13 (0.98–1.30) 0.93 (0.69–1.24)		0.394	1.00 (Ref)	0.97 (0.85–1.10)	0.99 (0.87–1.13)	0.935	1.00 (Ref)	1.05 (0.92–1.21)	1.05 (0.92–1.21) 1.14 (0.98–1.31)	0.083
(95% CI) Regional/distant metastatic N = 2187 1726 Age-adjusted OR 1.00	astatic 1726 1.00 (Ref)		371	<0.001	506 1.00 (Ref)	822 1.01 (0.88–1.17)	859 1.05 (0.91–1.21) 0.449	0.449	1487 1.00 (Ref)	323 0.97 (0.83–1.13)	377 1.31 (1.12–1.52)	0.003
(95% CI) Multivariable adjusted OR (95% CI)	1.00 (Ref)	1.00 (Ref) 1.21 (1.04–1.42) 1.10 (0.80–1.49)	1.10 (0.80–1.49)	0.048	1.00 (Ref)	0.97 (0.83–1.12)	0.97 (0.83–1.12) 0.95 (0.82–1.10) 0.515	0.515	1.00 (Ref)	0.92 (0.78–1.07)	0.92 (0.78–1.07) 1.17 (0.99–1.37)	0.172
Multivariable models-adjusted for age educational level, Ch diagnosis. Ref, Reference. ¹ Not adjusted for Charlson comorbidity index.	s-adjusted fo	r age educational le	svel, Charlson comc	orbidity in	ndex, serum	glucose, total chol	iarlson comorbidity index, serum glucose, total cholesterol, triglycerides, fasting status, time between measurement, and prostate cancer	s, fasting	j status, time	e between measure	ment, and prostate	Car

<5.60 5.60–6.90 PSA <4 versus 4.0–9.9($\mu g/L$) 971 152 N = 6126 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.06 (0.88–1.28) Multivariable adjusted 1.00 (Ref) 1.08 (0.89–1.30) 0R (95% Cl) PSA 4.0–9.9 ($\mu g/L$) (Reference level for outcome studied) N = 1164 781 PSA 4.0–9.9 ($\mu g/L$) (Reference level for outcome studied) N = 1164 781 PSA 10.0–20.0 versus 4.0–9.9($\mu g/L$) 781 781 PSA 10.0–20.0 versus 4.0–9.9($\mu g/L$) 781 781 PSA 10.0–20.0 versus 4.0–9.9($\mu g/L$) 781 781 PSA 10.0–20.0 versus 4.0–9.9($\mu g/L$) 781 781 PSA 20.0 versus 4.0–9.9($\mu g/L$) 1.00 (Ref) 1.05 (0.92–1.11) OR (95% Cl) 1.00 (Ref) 1.05 (0.92–1.11) OR (95% Cl) 1.00 (Ref) 1.05 (0.92–1.11) OR (95% Cl) 1.00 (Ref) 1.26 (1.35–1.11) OR (95% Cl) 1.00 (Ref) 1.28 (1.13–1.45) OR (95% Cl) 1.00 (Ref) 1.28 (1.13–1.45) OR (95% Cl) 1.00 (Ref) 1.28 (1.13–1.45)		>6.90 trend 1 41 (1.00-2.01) 0.081 43 (1.01-2.05) 0.071			P for		(P for
PSA <4 versus 4.0–9.9(μ g/L) 971 152 N = 6126 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.06 (0.88–1 Multivariable adjusted 1.00 (Ref) 1.06 (0.89–1 OR (95% Cl) PSA 4.0–9.9 (μ g/L) (Reference level for outcome studie N = 1164 5184 781 N = 1164 781 $N = 3219$ 0.0–20.0 versus 4.0–9.9(μ g/L) N = 3219 2.684 442 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.05 (0.92–1 Multivariable adjusted 1.00 (Ref) 1.05 (0.92–1 Multivariable adjusted 1.00 (Ref) 1.05 (0.92–1 N = 3351 2.655 595 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.41 (1.25–1 N = 3351 2.625 595 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.28 (1.13–1 OR (95% Cl) 1.00 (Ref) 1.28 (1.13–1 OR (95% Cl) 2.00 (Ref) 1.00 (Ref) 1.28 (1.13–1 OR (95% Cl) 2.00 (Ref) 1.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (85% Cl) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (85% Cl) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (85% Cl) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (85% Cl) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (85% Cl) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (Ref) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (Ref) 2.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (Ref) 2.00 (Ref) 1.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (Ref) 2.00 (Ref) 1.00 (Ref) 1.28 (1.13–1) N = 3351 0.0 (Ref) 2.00 (Ref) 2.0			d <5.18	5.18-6.19	>6.19 trend	<1.70	1.70-2.24	≥2.25 trend
Age-adjusted OR (95% CI) 1.00 (Ref) 1.06 (0.88–1 Multivariable adjusted 1.00 (Ref) 1.08 (0.89–1 OR (95% CI) 0R (95% CI) 1.00 (Ref) 1.08 (0.89–1 OR (95% CI) 0R (95% CI) 1.00 (Ref) 1.08 (0.89–1 PSA 4.0–9.9 (µg/L) Reference level for outcome studie 781 PSA 10.0–20.0 versus 4.0–9.9(µg/L) 781 781 PSA 10.0–20.0 versus 4.0–9.9(µg/L) 2684 442 Age-adjusted OR (95% CI) 1.00 (Ref) 0.98 (0.85–1 Multivariable adjusted 1.00 (Ref) 0.98 (0.85–1 OR (95% CI) 1.00 (Ref) 1.42 PSA >20.0 versus 4.0–9.9(µg/L) 781 781 PSA >20.0 versus 4.0–9.9(µg/L) 781 781 OR (95% CI) 1.00 (Ref) 1.41 (1.25–1 Multivariable adjusted 1.00 (Ref) 1.28 (1.13–1 OR (95% CI) 1.00 (Ref) 1.28 (1.13–1 OR (95% CI) 1.00 (Ref) 1.28 (1.13–1 OR (95% CI) 1.00 (Ref) 1.28 (1.13–1	(8) (0) (1) (0) (1) (1) (1) (1) (1) (1) (1) (1) (1) (1		329	448	387	816	188	160
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00 (Ref) 00 (Ref) 25 00 (Ref) 00 (Ref)	- 0 -		749	1253	1217	2205	507	507
25 20 (Ref) 00 (Ref)	>	.04 (0.80–1.35) 0.485 91 (0 0.450	5 1.00 (Ref)	1.09 (0.98–1.22) 1.05 (0.94–1.18)	1.03 (0.92–1.16) 0.754 0 98 (0 87–1 10) 0.576	1.00 (Ref) 1.00 (Ref)	1.05 (0.93–1.19) 1.04 (0.92–1.18)	1.13 (1.00–1.28) 0.044 1.11 (0.97–1.26) 0.119
(25 20 (Ref) 20 (Ref)		8)				(1001) 00: 1	1	(01)
N = 3351 2625 595 Age-adjusted OR (95% Cl) 1.00 (Ref) 1.41 (1.25-1 Multivariable adjusted 1.00 (Ref) 1.28 (1.13-1 OR (95% Cl) 0.00 (Ref) 1.28 (1.13-1								
Age-adjusted OR (95% Cl) 1.00 (Ref) 1.41 (1.25–1 Multivariable adjusted 1.00 (Ref) 1.28 (1.13–1 OR (95% Cl) Glascon 66 (Reference level for outcome studied)	131		789	1232	1330	2284	505	562
OR (95% CI) GR (95% CI) Glascon <6 (Reference lavel for nutrome studied)	~ ~	.41 (1.10-1.81) <0.001	11 1.00 (Ref)	0.97 (0.86–1.09)	1.00 (0.89–1.13) 0.889 0.01 (0.81–1.03) 0.153	1.00 (Ref) 1.00 (Ref)	1.01 (0.89–1.14) 0 08 /0 86–1.11)	1.22 (1.08–1.38) 0.004
Glaason <6 (Rafaranca laval for outcoma studiad)	-							
N = 6942 5859 896	187		1175	2637	2530	4841	1059	1042
Gleason 7 versus ≤6								
N = 4789 3913 727	149		1195	1795	1799	3314	733	742
CI) 1.00 (Ref)	<u></u>					1.00 (Ref)	1.02 (0.91–1.14)	
Multivariable adjusted 1.00 (Ref) 1.18 (1.05–1.33)	, -	.09 (0.84–1.40) 0.017	7 1.00 (Ref)	0.97 (0.87–1.07)	0.98 (0.88–1.08) 0.690	1.00 (Ref)	1.03 (0.92–1.17)	1.07 (0.95–1.20) 0.264
OR (95% CI)								
Gleason ≥8 versus ≤6								
1830	σ			869		1557	367	
CI) 1.00 (Ref)	<u></u>		-	1.06 (0.92–1.22)	1.14 (1.00–1.31)	1.00 (Ref)	1.10 (0.95–1.28)	1.25 (1.09–1.45)
Multivariable adjusted 1.00 (Ref) 1.24 (1.07–1.44) OR (95% CI)	<u> </u>	.37 (1.02–1.87) 0.001	1.00 (Ref)	1.05 (0.91–1.21)	1.11 (0.96–1.28) 0.156	1.00 (Ref)	1.06 (0.92–1.24)	1.17 (1.01–1.37) 0.040
T1–T2 (Reference level for outcome studied)								
N = 11087 9256 1525	306		2814	4197	4076	7725	1681	1681
T3-T4 versus T1-T2								
2226	-		656	1040	1119	1888	455	
1.00 (Ref)	-		-		1.04 (0.94–1.17)	1.00 (Ref)	1.12 (0.99–1.26)	1.18 (1.05–1.33)
Multivariable adjusted OR 1.00 (Ref) 1.12 (0.99–1.26) (95% Cl)	<u> </u>	.20 (0.96–1.52) 0.022	: 1.00 (Ref)	0.95 (0.85–1.06)	0.98 (0.88–1.10) 0.860	1.00 (Ref)	1.08 (0.96–1.22)	1.12 (0.99–1.27) 0.046

4, 30-32]. Conversely, several other studies reported a protective effect of hyperglycemia or diabetes against higher grade or more advanced PCa [33-38]. The observed positive association reported by our study may be explained by several underlying biological mechanisms. For several decades, glucose has been documented as an important source of energy for rapid tumor cell proliferation [39, 40]. Evidence from clinical and genetic studies have also linked the hyperglycemic environment to carcinogenic processes such as apoptosis, oxidative stress, DNA damage, and chronic inflammation, which may drive the aggressiveness and progression of cancer (Fig. 1) [11-14, 40, 41]. For instance, one mice study found that translation of the glycolytic enzyme hexokinase 2 (HK2) was increased in PCa cells due to loss of Pten and p53, which help to prevent cells from growing uncontrollably [11]. GLUT12, an important protein in the glycolytic pathway, was also observed to be highly expressed by PCacells and may potentially help to facilitate the high energy needs of tumor cells [12]. Other metabolic changes such as hyperinsulinemia, increased level of bioavailable IGF 1, and increased production of advanced glycation end products (AGE), which occur in response to a hyperglycemic environment, have also been linked to increased proliferation of cancer cells and poorer cancer outcomes [42, 43].

Furthermore, the observed positive association between glucose and more aggressive or advanced PCa persisted even after correcting for the potential confounding effects of BMI. Unlike most of the observational studies which examined the link between diabetes and PCa outcomes, we focused on serum glucose which has been suggested to be more sensitive than diabetes for accurately determining the relationship between glucose regulation and disease outcomes [44]. Nevertheless, in our study, we lacked information on duration of glucose aberrations. Prolonged glucose impairment has been reported to cause destruction of the Leydig cells, thereby, resulting in lower testosterone levels and potentially lowering the risk of worse outcomes [40, 45–48]. Nonetheless, evidence linking

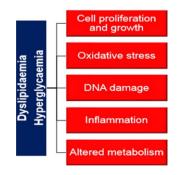


Figure 1. Schematic representation of carcinogenic processes suggested to be associated with hyperglycemia and dyslipidemia.

lower testosterone to reduced risk of worse PCa outcomes remains conflicting [46–48].

Similar to our study, one recent study reported higher PSA levels among prediabetic men [49]. In line with our findings, several studies also reported that diabetes or glucose in the diabetic range was associated with lower PSA levels [50-54]. Overall, our findings suggest that the effect of glucose levels in the prediabetic phase on PSA levels may be different from the diabetic phase. The potential mechanism through which prediabetes or impaired glucose tolerance may affect PSA levels is unknown. It is possible that in the prediabetic spectrum, the hormonal milieu characterized by hyperinsulinemia or higher IGF-1 activity may play a role [41]. With regard to men with glucose in the diabetic range, the underlying mechanisms which may partly explain the observed lower PSA levels involve reduction in testosterone levels and insulin-growth factor-1 bioactivity, resulting from hypoinsulinemia in the long term, and the use of metformin, a potential antineoplastic treatment [40, 41, 55-57]. Additionally, obesity, which is common among men with abnormal glucose levels, may also have contributed to the observed lower PSA levels due to PSA hemodilution [40, 52, 58]. However, in our subgroup analyses involving men with information on BMI, adjusting for BMI had minimal or no effect on the multivariate model.

Interestingly, we also found that although among men with glucose in the diabetic range (>6.9 mmol/L) had lower PSA levels, they had a greater odds of high-grade or more advanced PCa. This finding may be partly due to the fact that men with PSA <4 μ g/L are not likely to be biopsied and may therefore be diagnosed in the symptomatic stage when the cancer is more aggressive or advanced. These results are also in line with other studies indicating that men with low-PSA producing cancers tend to develop very poorly differentiated or highly tumorigenic castration-resistant PCa cells [59–61].

Total cholesterol and triglycerides

Four prospective studies found positive associations between total cholesterol and higher grade or more advanced PCa [5, 62–65]. Comparable to our study, the effect was positive, but statistically significant in two of these studies [5, 65]. Upon adjustment for age, we observed that the statistical significant ORs disappeared, which is consistent with the well-established view that age is a shared risk factor for both hypercholesterolemia and cancer [45, 66, 67].

Previously, we provided evidence supporting a potential role for lipid metabolism in PCa development [16, 17]. Here, we provided further evidence showing that triglycerides may influence the aggressiveness and severity of PCa. Except for one small Japanese study, most epidemiological studies did not report a link between triglycerides and PCa prognostic outcomes [2]. However, experimental studies using in vitro models, have shown that triglyceride-rich remnant like particles induce carcinogenesis by upregulating cell signaling pathways, such as the MEK/ERK and Akt pathways, involved in controlling cell growth and proliferation, apoptosis, cell cycle arrest, and lipid biosynthesis (Fig 1) [7, 8, 68].

Consistent with results from previous studies, total cholesterol did not appear to have an influence on PSA levels [69, 70]. In contrast to our findings, one study including 6774 Chinese men, reported an inverse association between triglycerides and PSA levels [71]. Two other studies, however, did not find any association between triglycerides and PSA [69, 70]. Further studies are needed to assess the relation between these lipids and PSA levels in men with PCa, as most studies to date involve healthy men from a relatively low-risk Asian population.

Strengths and limitations

This is one of the largest prospective studies to investigate the relation between glucose and lipid metabolism and clinicopathological characteristics of PCa. Our study population was also selected from an almost complete populationbased PCa register. Another strength of our study is the use of prediagnostic exposure measurements, which were all measured at a central laboratory. This enabled us to assess relations between the exposures and outcomes temporally and minimized the potential influence of reverse causation. Moreover, the results of our sensitivity analyses did not indicate any evidence of reverse causation.

PCa is a slow-progressing cancer [72] with a long induction period estimated at 6-13 years by some studies [73]. In this study, the relatively long time of 16.7 years between exposure measurement and PCa diagnosis, which may reflect the induction period between the exposures and PCa, can also be considered as a strength. However, we only used a single baseline exposure measurement, precluding consideration of the effect of longitudinal changes in exposure level over time [33, 37]. Further studies using repeated measurements are therefore warranted to account for the potential effect of this limitation. Our study also lacks information on participants' diabetic status and history of use of metformin and/or statin, so that exposure misclassification may have occurred. However, any such misclassification would be nondifferential since we used prediagnostic baseline exposure measurements. We did not have information on life-style-related factors such as smoking status and dietary history. Nonetheless, existing evidence does not appear to substantiate a link between the afore-mentioned lifestyle risk factors and poorer PCa

outcomes [1]. Based on the results of our sensitivity analyses, the lack of information on fasting status for some men did not appear to influence our results. Finally, we lack follow-up information on the clinicopathological characteristics of PCa such as PSA, and therefore, could not evaluate the association between the studied serum biomarkers and PCa biochemical recurrence or PCa progression.

Conclusion

Our findings suggest that the potential influence of glucose on PSA levels may vary as one progress from the prediabetic phase to the diabetic phase. Interestingly, though men in both the intermediate and highest glucose subgroups had poorer PCa outcomes. These findings may have important clinical implications for PCa detection and treatment, particularly among men with high glucose and low PSA levels, as previous evidence have shown that low PSA level is also concomitant with poorer PCa survival [61]. Overall, our findings further support a potential role for the lipid and glucose metabolism in prostate carcinogenesis.

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Conflict of Interests

None declared.

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Supporting Information

Additional supporting information may be found in the online version of this article:

Figure S1. Schematic representation of study population. **Table S1.** Odds ratios (OR) and 95% CI for prostate cancer risk categories by baseline levels of serum glucose, total cholesterol, and triglycerides for men with measurements more than 2 years prior to diagnosis.

Table S2. Odds ratio and 95% CI for the association between serum lipids (total cholesterol and triglycerides) and glucose levels and PSA levels and Gleason score.