Age and racial differences among PSA-detected (AJCC stage T1cN0M0) prostate cancer in the U.S.: a population-based study of 70,345 men

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Hong Zhang, Department of Radiation Oncology, University of Rochester Medical Center, 601 Elmwood Avenue Box 647, Rochester, NY 14642, USA e-mail: hong_zhang@ urmc.rochester.edu **Purpose**: Few studies have evaluated the risk profile of prostate-specific antigen (PSA)detected T1cN0M0 prostate cancer, defined as tumors diagnosed by needle biopsy because of elevated PSA levels without other clinical signs of disease. However, some men with stageT1cN0M0 prostate cancer may have high-risk disease (HRD), thus experiencing inferior outcomes as predicted by a risk group stratification model.

Methods: We identified men diagnosed with stageT1cN0M0 prostate cancer from 2004 to 2008 reported to the surveillance, epidemiology, and end results (SEER) program. Multivariate logistic regression was used to model the probability of intermediate-risk-disease (IRD) (PSA \geq 10 ng/ml but <20 ng/ml and/or GS 7), and high-risk-disease (HDR) (PSA \geq 20 ng/ml, and/or GS \geq 8), relative to low-risk disease (LRD) (PSA < 10 ng/ml and GS \leq 6), adjusting for age, race, marital status, median household income, and area of residence.

Results: A total of 70,345 men with PSA-detected T1cN0M0 prostate cancer were identified. Of these, 47.6, 35.9, and 16.5% presented with low-, intermediate-, and high-risk disease, respectively. At baseline (50 years of age), risk was higher for black men than for whites for HRD (OR 3.31, 95% CI 2.85–3.84). The ORs for age (per year) for HRD relative to LRD were 1.09 (95% CI 1.09–1.10) for white men, and as 1.06 (95% CI 1.05–1.07) for black men. Further, among a subgroup of men with low PSA (<10 ng/ml)T1cN0M0 prostate cancer, risk was also higher for black man than for white men at baseline (50 years of age) (OR 2.70, 95% CI 2.09–3.48). The ORs for age (per year) for HRD relative to LRD were 1.09 (95% CI 1.09–1.10) for white men, and as 1.06 (95% CI 1.05–1.07) for black men.

Conclusion: A substantial proportion of men with PSA-detected prostate cancer as reported to the SEER program had HRD. Black race and older age were associated with a greater likelihood of HRD.

Keywords: prostate cancer, race, age, population-study, SEER, screen-detected

INTRODUCTION

Prostate cancer is the most common malignancy in U.S. men. In 2013 alone, an estimated 241,000 new cancer cases will be diagnosed and 28,000 deaths will be attributed to prostate cancer (1). Current screening methods for prostate cancer include prostate-specific antigen (PSA) testing and digital rectal examination, although benefits of the former remain controversial (2, 3). The concern is that early detection and treatment of clinically insignificant prostate cancer may cause unnecessary side effects without added benefit.

Ever since FDA approval of PSA as a screening tool, many men have had a prostate biopsy because of an elevated PSA, despite a normal digital rectal examination. Prostate cancer diagnosed in this setting is classified as stage T1c disease based on the American Joint Commission on Cancer (AJCC) (4). Early studies have shown that stage T1c disease is heterogeneous in its pathological features. After a retrospective review of 257 patients with stage T1c prostate cancer who underwent prostatectomy and nodal dissection from 1987 to 1991, Lerner et al. (5) from the Mayo Clinic reported that 45% of patients had non-organ confined pT3 disease and 4% had node positive disease. Of 240 men with stage T1c disease who underwent radical prostatectomy at Johns Hopkins from 1994 to 1996, 28% had extracapsular extension, seminal vesicle, or lymph node involvement (6). From 1988 to 1998, 638 men with stage T1c prostate cancer underwent radical prostatectomy at Washington University, and 30% had non-organ confined pT3 or node positive disease (7). Many studies have shown that adverse pathological features such as pT3 disease (extracapsular extension and seminal vesicle involvement), high Gleason score (GS) and positive surgical margins increased the risk of disease

recurrence resulting in inferior outcomes (8-17). These studies suggested that some men with stage T1c disease might have high-risk prostate cancer. However, current PSA screening protocols are not able to identify clinically significant disease in this cohort.

The purpose of this study is to provide a contemporary profile of stage T1c prostate cancer based on demographic features and a risk stratification scheme developed by D'Amico et al. (18, 19) that includes stage, GS, and PSA level. This risk stratification has been validated and widely used, including by the National Comprehensive Cancer Network (NCCN). We analyzed demographic and tumor characteristics of over 70,000 men who, according to data reported to the Surveillance, Epidemiology, and End Results (SEER) program (2004–2008), were diagnosed with prostate cancer based on an elevated PSA level and without other clinical signs of disease (stage T1cN0M0). We report the probability of high-risk prostate cancer in these patients adjusting for characteristics such as age, race, marital status, median household income, and area of residence, while taking into account pre-biopsy PSA levels. These findings may provide important information in our efforts to develop more effective prostate cancer screening tools.

MATERIALS AND METHODS

PATIENT DATABASE

Men diagnosed with AJCC stage T1cN0M0 prostate adenocarcinoma at age \geq 18 years between 2004 and 2008 and reported to the SEER 17 Registries were identified. Year 2004 was chosen as the start year, since it was February 2004 that SEER initiated collection of detailed T, N, and M staging information. "Death certificate only" and "autopsy only" cases were excluded. A total of 78,367 cases were identified with stage T1cN0M0 disease. A total of 70,345 cases had PSA or GS available for analysis.

The youngest man in the cohort with stage T1cN0M0 disease was 37 years of age. We identified 262,172 men \geq 37 years of age with all stages of prostate cancer between 2004 and 2008. The SEER program provided the number of all men age \geq 37 years in SEER catchment area based on 2000 U.S. population estimates.

Based on PSA levels and GS, we divided cases into the three risk groups as described by D'Amico et al. (18). Low-risk disease (LRD) was defined as PSA <10 ng/ml and GS \leq 6; intermediate-risk disease (IRD) as PSA \geq 10 and <20 ng/ml and/or GS 7; and high-risk disease (HRD) as PSA \geq 20 ng/ml and/or GS \geq 8.

Patient socioeconomic status was evaluated using "median household income" and "rural-urban continuum code" provided in the SEER program, as in prior studies (20). Median household income is an aggregate measure based on county attributes derived from the 2000 U.S. census. Rural-urban continuum code, which provides information about potential accessibility to cancer care, allows classification of counties by population size, degree of urbanization, and proximity to metropolitan areas. We grouped patients into three categories by area of residence: within metropolitan areas, adjacent to metropolitan areas, and not adjacent to metropolitan areas. Marital status at diagnosis was classified as follows: never-married (single), married, and others (widowed, divorced, and separated).

STATISTICAL ANALYSIS

The means of continuous variables (e.g., age and median household income) were compared between groups using *t*-tests with unequal variances. Chi-square tests were used to assess differences in categorical variables (e.g., race and gender) between groups. Multivariate logistic regression analyses were conducted to model the probability of developing IRD and HRD. The list of potential predictors to be included in the models was age, race, marital status, median household income, area of residence, and all possible first-order interactions. The analyses treated "age" as continuous variable and set 50 year of age as baseline. To select predictors significantly associated with disease risk, the data set was randomly partitioned into a training and validation data set of equal sizes (50% of the original data set each). We first ran a backward model selection procedure on the training data set to identify candidate predictors potentially associated with disease risk. Once this training step was completed, we fitted the identified model to the validation data set; predictors with p-values smaller than 0.05 were considered significant and included in the final models. This analysis was performed for the IRD and HRD separately. The final models were also used to model the probability of developing IRD and HRD in a subgroup analysis for patients with PSA <10 ng/ml stage T1cN0M0 prostate cancer. In all of our statistical analyses, tests were two-sided and the significance level (probability of type-1 error) was set at 0.05. Analyses were conducted using the SAS statistical package, version 9.13 (SAS Institute, Cary, NC, USA).

RESULTS

The age distribution of men \geq 37 years of age in the SEER catchment areas are shown in **Table 1**. While 11.8% of men \geq 37 years of age in the SEER catchment area were \geq 75 years of age, 24.2% men with prostate cancer of all stages and 26.1% men with stage T1cN0M0 prostate cancer were \geq 75 years of age. Among all men diagnosed with prostate cancer (n = 262,172) in the SEER program, 78,367 (29.9%) had stage T1cN0M0 disease. The percentage changed with age, 15.8, 24.6, 35.2, and 32.2% for age groups of 37–49, 50–64, 65–74, and \geq 75 years, respectively (**Table 1**).

Table 1 | Age distribution of men in SEER catchment areas, and men with prostate cancer of all stages and stage T1cN0M0 specifically.

Age	No. of men in SEER areas ^a	No. of men with prostate cancer			
		All stages ^b	T1cN0M0 ^b	Percent (T1cN0M0/ all stage)	
37–49	14,983,548	1,550	244	15.8%	
50–64	12,929,972	19,646	4,826	24.6%	
65–74	4,530,521	18,520	6,510	35.2%	
≥75	4,329,573	12,718	4,093	32.2%	

^a Provided by SEER program using the 2000 U.S. standard population estimates. ^bAverage annual number of men with all stage prostate cancer and with stage T1cN0M0 prostate cancer are defined as crude rates reported by SEER between 2004 and 2008 divided by 5. A total of 70,345 men with stage T1cN0M0 prostate cancer had GS and PSA information available and these men were evaluated (median age 69 years, range 37–105) (**Table 2**). There were 11,600 men (16.5%) with HRD. Men with HRD were significantly older (median age 72, 70, and 67 for HRD, IRD, and LRD respectively, p < 0.01); more likely to be black (19.4 and 15.6% for black and white respectively, p < 0.01); less likely to be married (16.7 and 15.5% for never-married and married respectively); and had a lower median household income (17.7, 16.6, and 15.6% for first, second, and third tertile respectively, p < 0.01).

Based on age and PSA level, **Table 3** shows the distribution of GS among 70,345 men with stage T1cN0M0 disease. For example, a 72-year-old man with a PSA of 8 ng/ml stage T1cN0M0 prostate cancer has a 7.2% chance of having GS \geq 8 HRD. For an 82-year-old man with the same PSA level, the chance of HRD is 16.5%.

Older men were more likely to have higher PSA levels at the time of diagnosis (**Figure 1A**, p < 0.01). The percentage of men with PSA ≥ 20 ng/ml stage T1cN0M0 prostate cancer increased with age (5.3, 6.4, 7.6, and 13.3% for age groups 37– 49, 50–64, 65–74, and ≥ 75 , respectively). Furthermore, the percentage of men with higher GS increased with age (**Figure 1B**, p < 0.01). There were 3.6, 6.1, 9.1, and 17.5% men with GS ≥ 8 stage T1cN0M0 prostate cancer for age groups 37–49, 50– 64, 65–74, and ≥ 75 , respectively. Black men were more likely to have higher PSA levels than white men at diagnosis (**Figure 2A**, p < 0.01). For example, 10.6% of black men and 7.5% of white men had PSA of ≥ 20 ng/ml. There was a small but significant

Table 2 Characteristics of 70,345 men with low	-, intermediate-, and high-risk prostate cancer.
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	Low risk ^a	Intermediate risk ^a	High risk ^a	<i>p</i> -Value
Number of patients	33,472 (47.6%)	25,273 (35.9%)	11,600 (16.5%)	
Age, year (median, range)	67 (37–105)	70 (38–98)	72 (38–98)	< 0.01
<50	720 (68.6%)	248 (23.6%)	81 (7.7%)	
50–64	12,819 (58.6%)	6,597 (30.2%)	2,449 (11.2%)	
65–74	14,418 (48.6%)	10,869 (36.6%)	4,393 (14.8%)	
≥75	5,515 (31.1%)	7,559 (42.6%)	4,677 (26.3%)	
Race				< 0.01
White	26,465 (49.2%)	18,933 (35.2%)	8,366 (15.6%)	
Black	4,383 (41.5%)	4,118 (39.0%)	2,051 (19.4%)	
Other ^b	1,621 (40.8%)	1,511 (38.1%)	839 (21.1%)	
Unknown	1,003 (48.7%)	711 (34.5%)	344 (16.7%)	
Marital status				< 0.01
Married	23,308 (49.1%)	16,801 (35.4%)	7,371 (15.5%)	
Never-married	3,094 (47.0%)	2,391 (36.3%)	1,099 (16.7%)	
Other ^c	3,830 (42.2%)	3,475 (38.3%)	1,772 (19.5%)	
Unknown	3,240 (45.0%)	2,606 (36.2%)	1,358 (18.9%)	
Median household income ^d				< 0.01
First tertile	8,795 (46.1%)	6,919 (36.2%)	3,374 (17.7%)	
Second tertile	10,613 (47.2%)	8,149 (36.2%)	3,728 (16.6%)	
Third tertile	14,061 (48.9%)	10,205 (35.5%)	4,498 (15.6%)	
Area of residence				< 0.01
Within metro	30,493 (47.8%)	22,948 (36.0%)	10,347 (16.2%)	
Adjacent to metro	1,807 (46.0%)	1,390 (35.4%)	731 (18.6%)	
Not adjacent to metro	1,138 (44.5%)	910 (35.6%)	510 (19.9%)	
Unknown	33 (47.1%)	25 (35.7%)	12 (17.1%)	
Median follow up in months (range)	26 (12–59)	25 (1–59)	25 (1–59)	
Year of diagnosis				p = 0.60
2004	6,176 (48.5%)	4,421 (34.7%)	2,146 (16.8%)	
2005	5,976 (47.8%)	4,425 (35.4%)	2,104 (16.8%)	
2006	7,296 (48.4%)	5,290 (35.1%)	2,489 (16.5%)	
2007	7,508 (47.2%)	5,855 (36.8%)	2,527 (15.9%)	
2008	6,516 (46.1%)	5,282 (37.4%)	2,334 (16.5%)	

^a Risk groups: low risk – PSA < 10 ng/ml and GS ≤6, intermediate risk – 10 ng/ml ≤ PSA < 20 ng/ml and/or GS = 7, high risk – PSA > 20 ng/ml, and/or GS ≥8 as defined by D'Amico (19).

^bOther: American Indian, Alaska Native, Asian, and Pacific Islander.

^cOther: widowed, divorced, and separated.

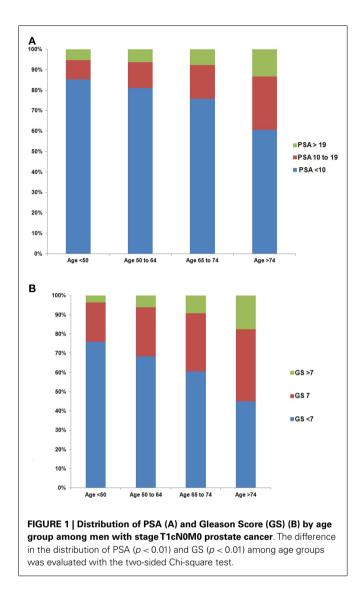
^dFirst tertile, < 42,190; second tertile, \$42,191–49,820; third tertile, \ge 49,821.

Age							Ā	PSA (ng/ml)							
		4			4-9.9			10–14.9			15–19.9			≥20	
<50	195 (84.8%)	33 (14.4%)	2 (0.9%)	525 (78.9%)	124 (18.6%)	15 (2.3%)	40 (49.4%)	34 (42.0%)	7 (8.6%)	10 (55.6%)	7 (38.9%)	1 (5.6%)	27 (48.2 <i>%</i>)	16 (28.6%)	13 (23.2%)
50–54	372 (81.9%)	75 (16.5%)	7 (1.5%)	1,541 (73.5%)	496 (23.7%)	58 (2.8%)	165 (58.7%)	92 (32.7%)	23 (8.2%)	45 (51.1%)	32 (36.4%)	10 (11.4%)	86 (37.7%)	100 (43.9%)	41 (18.0%)
55-59	647 (81.0%)	133 (16.6%)		3,771 (77.8%)	1,191 (23.0%)	221 (4.3%)	392 (58.5%)	219 (32.7%)	59 (8.8%)	100 (48.8%)	72 (35.1%)	33 (16.1%)	204 (44.2 <i>%</i>)	161 (34.8%)	97 (21.0%)
60-64	765 (75.4%)	198 (19.5%)	51 (5.0%)	5,723 (70.0%)	2,045 (25.0%)	411 (5.0%)	638 (59.6%)	344 (32.1%)	89 (8.3%)	201 (47.6%)	159 (37.7%)	62 (14.7%)	321 (44.7%)	253 (35.2%)	144 (10.1%)
65–69	806 (71.6%)	256 (22.7%)	64 (5.7%)	6,829 (66.3%)	2,868 (27.8%)	609 (5.9%)	893 (54.2%)	549 (33.3%)	206 (12.5%)	261 (49.7%)	185 (35.2%)	79 (15.0%)	389 (36.9%)	427 (40.6%)	237 (22.5%)
70–74	651 (66.6%)	246 (25.2%)	80 (8.2%)	6,136 (60.6%)	3,258 (32.2%)	733 (7.2%)	1,076 (54.2%)	663 (33.4%)	247 (12.4%)	352 (48.2%)	262 (35.8%)	117 (16.0%)	431 (35.8%)	452 (37.5%)	322 (26.7%)
75–79	365 (53.2%)		86 (12.5%)	3,769 (53.9%)	2,493 (35.6%)	736 (10.5%)	908 (46.3%)	708 (36.1%)	344 (17.6%)	274 (42.9%)	222 (34.7%)	143 (22.4%)	377 (33.4%)	441 (39.1%)	310 (27.5%)
80	335 (46.1%)	281 (38.7%)	110 (15.2%)	1,046 (44.3%)	925 (39.2%)	389 (16.5%)	481 (35.2%)	558 (40.8%)	328 (24.0%)	205 (31.4%)	269 (41.2%)	179 (27.4%)	248 (20.1%)	512 (41.5%)	474 (38.4%)
	LRD ^a	IRD ^b	HRD℃	LRD ^a	IRD ^b	HRD℃	IRD ^b	IRD ^b	HRD℃	IRD ^b	IRD ^b	HRD℃	HRD℃	HRD°	HRD℃
	9 ⊻I	7	8 ^I	9 VI	7	8 ^I	9 VI	7	8 ∧I	9 ∀I	7	8 ∧I	9 ∀I	7	8 ^I
							ß	Gleason score	۵)						

Table 3 | Distribution of low-, intermediate-, and high- risk T1cN0M0 prostate cancer according to age and PSA.

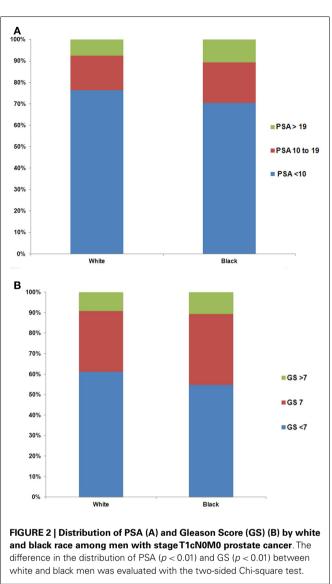
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difference in GS profile between black and white men (Figure 2B, p < 0.01).

Multivariate logistic regression analyses identified a significant interaction between age and race, indicating that risk increased faster with age among white men compared to blacks for both HRD and IRD relative to LRD (Table 4). At baseline (50 years of age), risk was higher among black men than among whites for HRD and IRD (OR 3.31 and 2.02, 95% CI 2.85-3.84, and 1.81-2.25, respectively). The ORs for age (per year) for HRD and IRD relative to LRD were estimated as 1.09 and 1.06 respectively (95% CI 1.09-1.10, 1.05–1.06) for white men, and as 1.06 and 1.04 respectively (95% CI 1.05-1.07, 1.03-1.04) for black men. Compared to baseline (50 years of age), the ORs at 75 years of age for HRD and IRD relative to LRD were estimated as 9.25 (95% CI 8.45-10.13) and 3.97 (95% CI 3.72-4.24) respectively for white men, and as 4.24 (95% CI 3.56-5.04) and 2.58 (95% CI 2.24-2.96) respectively for black men. Men in the "never-married" category compared with "married" men were more likely to have HRD and IRD (OR 1.35 and 1.22, 95% CI 1.25-1.46 and 1.15-1.29, respectively), but no



significant association was observed between median household income and disease risk.

As one may argue that the elevated PSA levels in the elderly patients with HRD may be a function of lead-time detection of younger individuals with lower PSA, we also analyzed age and racial effect in a subgroup of men with PSA <10 ng/ml stage T1cN0M0 prostate cancer (Table 5). We found that in this group of men with low PSA (<10 ng/ml), again, older men were more likely to have GS \geq 8 HRD (12.3, 6.6, 4.3, and 1.9% for age groups \geq 75, 65–74, 50–64, and 37–49 respectively). Black men were more likely to have GS \geq 8 HRD (7.6 and 6.7% for black and white men respectively). Multivariate logistic regression analyses for this cohort of men showed that risk increased faster with age among white men compared to black men for both HRD and IRD relative to LRD (Table 6). At baseline (50 years of age), risk was higher among black men than among whites for HRD and IRD (OR 2.70 and 1.94, 95% CI 2.09-3.48 and 1.70-2.20, respectively). The OR for age (per year) for patients with GS ≥ 8 HRD (relative to

	Inte	rmediate vs. low risl	(^a		High vs. low risk ^a	
	Odds ratio	95% Cl ^b	<i>p</i> -Value	Odds ratio	95% Cl ^b	<i>p</i> -Value
Age (per year) × race						
White	1.06	(1.05, 1.06)	< 0.01	1.09	(1.09, 1.10)	< 0.01
Black	1.04	(1.03, 1.04)	< 0.01	1.06	(1.05, 1.07)	< 0.01
Other ^c	1.06	(1.04, 1.07)	0.86	1.08	(1.07, 1.11)	0.27
Race			< 0.01			< 0.01
White	_	_	_	_	_	-
Black	2.02	(1.81, 2.25)	< 0.01	3.31	(2.85, 3.84)	< 0.01
Other ^c	1.30	(1.06, 1.60)	0.01	1.58	(1.40, 2.46)	< 0.01
Marital status			< 0.01			< 0.01
Married	_	_	_	_	_	-
Never-married	1.22	(1.15, 1.29)	< 0.01	1.35	(1.25, 1.46)	< 0.01
Others ^d	1.19	(1.13, 1.26)	< 0.01	1.34	(1.25, 1.43)	< 0.01
Median household income ^e			0.43			0.42
First tertile	-	-	-	-	-	-
Second tertile	0.98	(0.94, 1.03)	0.50	0.98	(0.92, 1.05)	0.56
Third tertile	0.97	(0.92, 1.02)	0.20	0.96	(0.90, 1.02)	0.20
Area of residence			0.66			< 0.01
Within metro	-	-	-	-	-	-
Adjacent to metro	1.00	(0.92, 1.09)	0.93	1.11	(1.00, 1.23)	0.06
Not adjacent to metro	1.05	(0.95, 1.16)	0.37	1.29	(1.13, 1.46)	< 0.01

Table 4 | Multivariate logistic regression analyses modeling the probabilities of patients with intermediate-risk disease vs. low-risk disease, and of patients with high-risk diseases vs. low-risk disease respectively.

^a Risk groups: low risk – PSA < 10 ng/ml and GS ≤6, intermediate risk – 10 ng/ml ≤ PSA <20 ng/ml and/or GS = 7, high risk – PSA ≥20 ng/ml and/or GS ≥8. ^b Cl, confidence interval.

^cOthers: American Indian, Alaska Native, Asian, and Pacific Islander.

^dOther: widowed, divorced, and separated.

^e First tertile, <\$ 42,190; second tertile, \$ 42, 191–49,820; third tertile, ≥\$ 49,821.

LRD) were 1.09 (95% CI 1.09–1.10) for white men and 1.06 (95% CI 1.05–1.07) for black men. Compared to baseline 50 years of age, the ORs for 75-year-old men for HRD and IRD relative to LRD were estimated as 8.51 (95% CI 7.36–9.86) and 3.25 (95% CI 3.01–4.51) respectively for white men, and as 4.09 (95% CI 3.05–5.50) and 2.09 (95% CI 1.78–2.46) respectively for black men.

DISCUSSION

To our knowledge, this is the largest population-based study focused only on PSA-detected (stage T1cN0M0) prostate cancer in the U.S. in the contemporary era of widespread PSA testing. A significant number of men (16.5%) in this cohort had HRD. We found that men of older age and black race were more likely to have HRD than younger and white men.

According to the U.S. census, 4.6% of the total U.S. male population in 2010 was \geq 75 years of age respectively (21). However, 40.3% of men with HRD were \geq 75 years of age. Further, 36.8% of men with PSA <10 ng/ml and GS \geq 8 stage T1cN0M0 prostate cancer were \geq 75 years of age. Our finding that older men were more likely to have HRD, albeit limited to PSA-detected stage T1cN0M0 disease in this survey, is consistent with recently published studies (22, 23). After evaluating all prostate cancer cases reported to the SEER program from 1998 to 2007, Scosyrev et al. (22) reported

that men \geq 75 years of age were more likely to present with either metastatic or locally advanced disease, and experienced the highest prostate cancer-specific mortality. Bechis et al. (23) reviewed the database of the Cancer of the Prostate Strategic Urologic Research Endeavor (CaPSURE) and reported that 26% of men age \geq 75 had HRD based on the Cancer of the Prostate Risk Assessment (CAPRA) score. In addition to clinical stage, PSA, and GS, CAPRA scores take into consideration patient age and percentage of biopsy cores involved with prostate cancer.

The specific indications for PSA testing and prostate biopsy could not be confirmed in our series, given the limitations of the SEER database. Nonetheless, all cases reported to the SEER program were diagnosed by a needle biopsy because of an elevated PSA level with no other clinical signs of disease (AJCC T1cN0M0) (4). The higher proportion of older men with HRD reported here cannot be explained by potential bias that older men with inherently higher PSA levels were more likely to have biopsy. Even among men with PSA <10 ng/ml stage T1cN0M0 disease, older men had significant risk of HRD. One of many possible explanations is that older men may harbor aggressive disease that is not reflected by PSA level. After pathological review of 211 autop-sied prostate glands from deceased men with no known prostate cancer at the time of death, Delongchamps et al. (24) reported older men had significantly larger tumors, higher GS, and were

Table 5 | Characteristics of 51,919 men with PSA < 10 ng/ml stage T1cN0M0 prostate cancer.

	Low risk ^a	Intermediate risk ^a	High risk ^a	<i>p</i> -Value
Number of patients	33472 (64.5%)	14857 (28.6%)	3590 (6.9%)	
Age, year (median, range)	67 (37–105)	70 (38–95)	72 (44–95)	<0.01
<50	720 (80.5%)	157 (17.6%)	17 (1.9%)	
50–64	12819 (72.3%)	4138 (23.3%)	766 (4.3%)	
65–74	14418 (64.0%)	6628 (29.4%)	1486 (6.6%)	
≥75	5515 (51.2%)	3934 (36.5%)	1321 (12.3%)	
Race				<0.01
White	26465 (65.4%)	11318 (28.0%)	2697 (6.7%)	
Black	4383 (59.9%)	2380 (32.5%)	560 (7.6%)	
Other ^b	1621 (61.2%)	778 (29.4%)	250 (9.4%)	
Unknown	1003 (68.4%)	381 (26.0%)	83 (5.7%)	
Marital status				<0.01
Married	23,308 (65.0%)	10,117 (28.2%)	2,444 (6.8%)	
Never-married	3,094 (65.9%)	1,302 (27.7%)	297 (6.3%)	
Other ^c	3,830 (61.5%)	1,891 (30.4%)	505 (8.1%)	
Unknown	3,240 (62.9%)	1,547 (30.4%)	344 (6.8%)	
Median household income ^d				<0.01
First tertile	8,795 (63.9%)	3,972 (28.9%)	996 (7.2%)	
Second tertile	10,613 (65.1%)	4,583 (28.1%)	1,108 (6.8%)	
Third tertile	14,061 (64.4%)	6,299 (28.8%)	1,486 (6.8%)	
Area of residence				<0.01
Within metro	30,493 (64.4%)	13,588 (28.7%)	3,263 (6.9%)	
Adjacent to metro	1,807 (64.5%)	790 (28.2%)	204 (7.3%)	
Not adjacent to metro	1,138 (65.7%)	471 (27.2%)	122 (7.0%)	
Year of diagnosis				
2004	6,176 (68.2%)	2,317 (25.6%)	563 (6.2%)	
2005	5,976 (66.3%)	2,421 (26.9%)	616 (6.8%)	
2006	7,296 (65.3%)	3,098 (27.7%)	779 (7.0%)	
2007	7,508 (62.8%)	3,597 (30.1%)	845 (7.1%)	
2008	6,516 (60.7%)	3,424 (31.9%)	787 (7.3%)	

 a Risk groups: low risk – PSA < 10 ng/ml and GS \leq 6, intermediate risk – PSA < 10 ng/ml and GS = 7, high risk – PSA < 10 ng/ml and GS \geq 8.

^bOther: American Indian, Alaska Native, Asian, and Pacific Islander.

°Other: widowed, divorced, and separated.

^dFirst tertile, <\$ 42,190; second tertile, \$ 42,191–49,820; third tertile, ≥\$ 49,821.

more likely to have extraprostatic extension or microscopic invasion of bladder neck (4). After a prospective review of 268 men with stage T1c prostate cancer who underwent radical prostatectomy at seven U.S. medical centers, Southwick et al. (25) noted age was one of the significant predictors of unfavorable pathological outcome, including extracapsular extension, seminal vesicle invasion, invasion of bladder neck/rectum, and lymph node involvement. Further study is needed to evaluate many confounding factors on observed age effect.

African Americans have the highest prostate cancer burden and mortality of any racial group (1). Our study showed that in the cohort of men with PSA screen-detected stage T1cN0M0 disease, African Americans had a higher likelihood of harboring HRD than whites, including cohort of men with PSA <10 ng/ml. Many socioeconomic and intrinsic factors may contribute to such a racial difference. But even among U.S. service men with equal access to care, racial difference in prostate cancer risk remains (26). In an early study among men with non-palpable prostate cancer (clinical stage T1c disease) who underwent prostatectomy, Sanchez-Ortiz et al. reported that African American men had higher GS and greater tumor volume (27). Therefore, prostate cancer in black men may be biologically different from whites. In fact, several genetic and biological mechanisms have been identified that may contribute to the aggressiveness of prostate cancer in African American men (28–32).

Our study was not designed to address the question of whether earlier detection would improve survival of men with HRD. Also to our knowledge, there have been no reported randomized studies that evaluate the outcome of early intervention vs. active surveillance for men found to have stage T1cN0M0 prostate cancer. There are two UK-based ongoing prostate cancer trials: the CAP (Comparison Arm for ProtecT) and ProtecT (Prostate Testing for Cancer and Treatment) trials (33) that may help address issues of screening and treatment, but the results will not be

	Inte	rmediate vs. low risl	(^a		High vs. low risk ^a	
	Odds ratio	95% Cl ^b	<i>p</i> -Value	Odds ratio	95% Cl ^b	<i>p</i> -Value
Age (per year) × race						
White	1.05	(1.04, 1.05)	<0.01	1.09	(1.09, 1.10)	<0.01
Black	1.04	(1.04, 1.05)	<0.01	1.06	(1.05, 1.07)	<0.01
Other ^c	1.04	(1.03, 1.06)	0.26	1.08	(1.06, 1.10)	0.26
Race			<0.01			<0.01
White	-	_	_	-	_	_
Black	1.94	(1.70, 2.20)	<0.01	2.70	(2.09, 3.48)	<0.01
Other ^c	1.27	(0.99, 1.63)	0.06	1.85	(1.17, 2.91)	<0.01
Marital status			<0.01			<0.01
Married	-	_	_	-	_	_
Never-married	1.07	(0.99, 1.15)	0.07	1.11	(0.97, 1.27)	0.12
Others ^d	1.09	(1.03, 1.16)	<0.01	1.18	(1.06, 1.31)	<0.01
Median household income ^e			0.09			0.45
First tertile	-	_	_	_	_	-
Second tertile	0.95	(0.90, 1.01)	0.12	0.95	(0.85, 1.05)	0.29
Third tertile	1.01	(0.95, 1.07)	0.82	0.99	(0.90, 1.10)	0.90
Area of residence			0.19			0.84
Within metro	-	_	_	_	_	-
Adjacent to metro	0.95	(0.86, 1.04)	0.27	0.98	(0.83, 1.67)	0.84
Not adjacent to metro	0.90	(0.80, 1.02)	0.11	0.94	(0.75, 1.17)	0.57

Table 6 | Multivariate logistic regression analyses modeling the probabilities of patients with PSA <10 ng/ml intermediate-risk disease vs. low-risk disease, and of patients with PSA <10 ng/ml high-risk diseases vs. low-risk disease respectively.

^aRisk groups: low risk – PSA < 10 ng/ml and GS ≤6, intermediate risk – PSA < 10 ng/ml and GS = 7, high risk – PSA < 10 ng/ml and GS ≥8.

^bCl, confidence interval.

^cOthers: American Indian, Alaska Native, Asian, and Pacific Islander.

^dOther: widowed, divorced, and separated.

^e First tertile, <\$ 42,190; second tertile, \$ 42,191–49,820; third tertile, ≥\$ 49,821.

available until 2016. Based on results from many published studies (34–36) including stage T1cN0M0 disease, it is conceivable that a subgroup of men may indeed benefit from early detection and treatment. The challenge at this time is to distinguish these patients from many men with clinically non-consequential disease.

Our study has several weaknesses due to the retrospective design and inherent deficiencies of data reported to the SEER program. We had no way to independently verify staging accuracy. SEER did not provide detailed information about biopsy templates/schemes. We were not able to analyze patient outcomes due to the short follow up in the cohort. Further, there was no information about patients' performance status, medical co-morbidities, voiding symptoms, or family history of prostate cancer in the SEER database; these factors might have influenced screening decisions. Eight percent (5,605/70,345) of men in this study had a PSA level <4 ng/ml. It is unclear why these men proceeded to biopsy, though likely explanations may include patient preference, family history of prostate cancer, or incorrect staging.

We used a risk stratification scheme developed by D'Amico et al. (18, 19) that has been validated and widely used. In addition to clinical stage, PSA, and GS, there are many other factors that may influence prostate cancer outcome. These include primary and secondary GS, tertiary GS (37-41), percentage of positive biopsies

(42–47), or presence of perineural invasion in the biopsy specimen (48). The SEER database does not provide such important information; therefore we limited our risk estimation based on clinical stage (in this case, stage T1cN0M0), PSA, and GS.

Despite the above limitations, our large population-based study shows that a substantial proportion of men with PSA-detected stage T1cN0M0 prostate cancer may have HRD in the contemporary era. Older and black men were more likely to have HRD than younger and white men. Analytic studies with independently verified staging information are needed to confirm these findings, and examine clinical outcomes in these men, especially those of older age and of black race.

AUTHOR CONTRIBUTIONS

Conception and design: Hong Zhang, Edward M. Messing, Lois B. Travis, and Yuhchyau Chen; Collection and assembly of data: Hong Zhang; Data analysis and interpretation: all authors; Manuscript writing: all authors; Final approval of manuscript: all authors.

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