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Platinum Priority – Prostate Cancer

Editorial by XXX on pp. x–y of this issue

Mortality Among Men with Advanced Prostate Cancer Excluded from the ProtecT Trial

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\textbf{Abstract}

\textbf{Background:} Early detection and treatment of asymptomatic men with advanced and high-risk prostate cancer (PCa) may improve survival rates.

\textbf{Objective:} To determine outcomes for men diagnosed with advanced PCa following prostate-specific antigen (PSA) testing who were excluded from the ProtecT randomised trial.

\textbf{Design, setting, and participants:} Mortality was compared for 492 men followed up for a median of 7.4 yr to a contemporaneous cohort of men from the UK Anglia Cancer Network (ACN) and with a matched subset from the ACN.

\textbf{Outcome measurements and statistical analysis:} PCa-specific and all-cause mortality were compared using Kaplan-Meier analysis and Cox’s proportional hazards regression.

\textbf{Results and limitations:} Of the 492 men excluded from the ProtecT cohort, 37 (8%) had metastases (N1, M0 = 5, M1 = 32) and 305 had locally advanced disease (62%). The median PSA was 17 µg/l. Treatments included radical prostatectomy (RP: n = 54; 11%), radiotherapy (RT: n = 245; 50%), androgen deprivation therapy (ADT; n = 122; 25%), other treatments (n = 11; 2%), and unknown (n = 60; 12%). There were 49 PCa-specific

\textsuperscript{1}These authors contributed equally to this work.

\textsuperscript{*}The members of the ProtecT study group are listed in the Supplementary material.

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1. Introduction

Population-based prostate specific antigen (PSA) screening remains controversial [1]. Although screening in the European Randomised Study of Screening for Prostate Cancer (ERSPC) detected high numbers of prostate cancers (PCAs) and lower mortality from that disease, the majority of cancers were indolent, leading to over-detection and overtreatment [2,3]. The Prostate, Lung and Ovarian cancer screening study (PLOCO) reported no survival benefit after 11.5 yr of follow-up, but there was widespread contamination in the control arm with previous PSA testing (up to 90%) [2,3].

There is uncertainty regarding the effectiveness of treatments for PSA-detected clinically localised PCAs. The Prostate Cancer Intervention Versus Observation Trial (PIVOT) reported no survival benefit after 12 yr of follow-up among men with mainly low-risk disease treated with surgery or observation, although there was high all-cause mortality in both arms, suggesting that men with major comorbidities were included [4]. No randomised trials have compared different radical treatments for men with advanced [5,6] or high-risk disease, and retrospective studies have reported conflicting results [7–10]. There is uncertainty regarding outcomes among men with higher-risk PCAs detected via PSA screening, although a subgroup analysis of PIVOT suggested benefit in favour of radical treatment for intermediate- or high-risk disease [4].

Details of the ProtecT trial are reported elsewhere [11–15]. Men with metastatic or locally advanced disease (cT3–4) and/or PSA ≥ 20 μg/l were excluded from ProtecT, along with men considered by local urologists to be unsuitable for the trial because of their clinical features. These men excluded from the ProtecT randomised trial but diagnosed contemporaneously provide a unique opportunity to assess the outcomes of advanced and high-risk disease at diagnosis in a population with very low rates of opportunistic PSA screening (8–13%) [12,16].

Here we present survival data for these men in comparison to data for a contemporaneous cohort from the UK Anglia Cancer Network (ACN), which has generally low rates of PSA testing, and with a matched ACN cohort with similar disease features.

2. Patients and methods

2.1. Case population

The ProtecT trial compares active monitoring, conformal external-beam radical radiotherapy (RT) with or without androgen deprivation therapy (ADT) and radical prostatectomy (RP) treatments for PSA-detected clinically localised PCA [12]. Between 2001 and 2009 there were 82 429 asymptomatic men aged 50 and 69 yr who underwent PSA testing, and those with PSA ≥ 3 μg/l proceeded to biopsy. Participants with initial PSA ≥ 20 μg/l or found to have locally advanced (T3–4) PCAs or distant disease (N1 or M1) were ineligible and referred for standard care. The majority had locally advanced PCAs; a small proportion (5%) were classed as at high risk of having non-organ-confined disease and were felt to be unsuitable for randomisation. In total, 513 men (PSA ≥ 20 μg/l, or locally advanced cT3–4 PCAs, or Gleason ≥ 8, or N1/M1 disease) were excluded from ProtecT (Table 1). These men form the ProtecT advanced cases cohort reported here.

Information on treatment and survival was obtained during annual ProtecT follow-up and checked using the English National Cancer Online Registration Environment database in the Eastern Office of the National Cancer Registration Service (NCRS-E) [15]. Cause of death was determined by review of certification by two independent clinicians blinded to study group and treatment.

2.2. Comparison population

Comparison patients (controls) were identified by the NCRS-E by interrogation of the Anglia Cancer Network (ACN) [10] for a contemporaneous cohort of men with comparable age and year of diagnosis and similarly advanced and high-risk disease features: PSA ≥ 20 ng/ml, locally advanced disease (cT3–4), Gleason score ≥ 8, or N1/M1 disease. The ACN cases were judged to be a suitable comparative cohort because of low rates of PSA testing (10–13%) in the ACN population [12,17] (Supplementary material).

The ProtecT trial was approved by the East Midlands Ethics Committee (Derby, UK; record number 01/4/025).

2.3. Statistical analysis

We used the χ² test for heterogeneity to assess baseline differences between cases and controls. The primary analysis compared risk of death from PCAs and all causes between ProtecT cases and ACN controls with clinically detected PCAs. Cases and controls were matched 1:1 according to age, year of diagnosis, PSA, Gleason score, and clinical stage. Survival estimates were carried out using Kaplan-Meier methods, with group differences (unmatched and matched) expressed as the hazard ratio (HR)
Table 1 – Demographic and clinicopathologic data for the study cohort

<table>
<thead>
<tr>
<th>Variable</th>
<th>Unmatched</th>
<th>Matched</th>
<th>p value</th>
<th>Unmatched</th>
<th>Matched</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Patients (n)</td>
<td>492</td>
<td>3978</td>
<td>1</td>
<td>401</td>
<td>401</td>
<td>1</td>
</tr>
<tr>
<td>Year of diagnosis, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1999–2003</td>
<td>178 (36)</td>
<td>1109 (28)</td>
<td>&lt;0.0001</td>
<td>151 (38)</td>
<td>151 (38)</td>
<td>1</td>
</tr>
<tr>
<td>2004–2006</td>
<td>191 (39)</td>
<td>1117 (28)</td>
<td></td>
<td>157 (39)</td>
<td>157 (39)</td>
<td></td>
</tr>
<tr>
<td>2007–2010</td>
<td>123 (25)</td>
<td>1752 (44)</td>
<td></td>
<td>93 (23)</td>
<td>93 (23)</td>
<td></td>
</tr>
<tr>
<td>Age band, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;50 yr</td>
<td>102 (21)</td>
<td>567 (14)</td>
<td>&lt;0.0002</td>
<td>83 (21)</td>
<td>81 (20)</td>
<td>0.86</td>
</tr>
<tr>
<td>50–72 yr</td>
<td>390 (79)</td>
<td>3411 (86)</td>
<td></td>
<td>318 (79)</td>
<td>320 (80)</td>
<td></td>
</tr>
<tr>
<td>Serum PSA, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;10 ng/ml</td>
<td>160 (33)</td>
<td>728 (18)</td>
<td>&lt;0.0001</td>
<td>149 (37)</td>
<td>144 (36)</td>
<td>0.48</td>
</tr>
<tr>
<td>10–20 ng/ml</td>
<td>116 (24)</td>
<td>752 (19)</td>
<td></td>
<td>112 (28)</td>
<td>117 (29)</td>
<td></td>
</tr>
<tr>
<td>20–50 ng/ml</td>
<td>141 (28)</td>
<td>1086 (27)</td>
<td></td>
<td>90 (22)</td>
<td>75 (19)</td>
<td></td>
</tr>
<tr>
<td>50–100 ng/ml</td>
<td>49 (10)</td>
<td>462 (12)</td>
<td></td>
<td>24 (6)</td>
<td>30 (7)</td>
<td></td>
</tr>
<tr>
<td>&gt;100 ng/ml</td>
<td>26 (5)</td>
<td>769 (19)</td>
<td></td>
<td>26 (6)</td>
<td>35 (9)</td>
<td></td>
</tr>
<tr>
<td>Unknown</td>
<td>0 (0)</td>
<td>181 (5)</td>
<td></td>
<td>0 (0)</td>
<td>0 (0)</td>
<td></td>
</tr>
<tr>
<td>Mean PSA, ng/ml (median)</td>
<td>32.6 (16.7)</td>
<td>201.1 (26.5)</td>
<td></td>
<td>31.7 (14)</td>
<td>217.2 (13)</td>
<td></td>
</tr>
<tr>
<td>Gleason score, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>112 (23)</td>
<td>473 (12)</td>
<td>93 (23)</td>
<td>92 (23)</td>
<td>1</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>259 (53)</td>
<td>1300 (33)</td>
<td>222 (53)</td>
<td>223 (55)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>115 (23)</td>
<td>1654 (42)</td>
<td>86 (21)</td>
<td>86 (21)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Unknown</td>
<td>6 (10)</td>
<td>551 (14)</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Mean Gleason score (median)</td>
<td>7.1 (7)</td>
<td>7.6 (7)</td>
<td>7.1 (7)</td>
<td>7.6 (7)</td>
<td>0.18</td>
<td></td>
</tr>
<tr>
<td>Clinical stage, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>T1</td>
<td>17 (4)</td>
<td>989 (25)</td>
<td>16 (4)</td>
<td>29 (7)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>T2</td>
<td>42 (8)</td>
<td>750 (19)</td>
<td>42 (10)</td>
<td>29 (7)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>T3</td>
<td>305 (62)</td>
<td>1063 (27)</td>
<td>301 (75)</td>
<td>298 (74)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>T4</td>
<td>5 (10)</td>
<td>44 (1)</td>
<td>4 (1)</td>
<td>5 (1)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>M1 or N1</td>
<td>37 (8)</td>
<td>1132 (28)</td>
<td>37 (9)</td>
<td>40 (10)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>T stage unknown</td>
<td>86 (18)</td>
<td>0 (0)</td>
<td>1 (1)</td>
<td>0 (0)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Follow-up (yr)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Mean</td>
<td>7.5</td>
<td>5.5</td>
<td>7.7</td>
<td>7.5</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Median</td>
<td>7.4</td>
<td>5</td>
<td>7.6</td>
<td>7.6</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Interquartile range</td>
<td>5.5–9.7</td>
<td>3.1–7.8</td>
<td>5.5–9.8</td>
<td>5.1–9.8</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

ACN = Anglia Cancer Network; PSA = prostate-specific antigen.

* p value for χ² test for heterogeneity between unmatched and matched ProtecT advanced cases and ACN controls.

with 95% confidence interval (CI) and compared using the log-rank test. Cox proportional hazards regression models (univariable and multivariable) were also fitted to estimate survival for the unmatched ProtecT cases and ACN controls adjusted for the above variables, with results expressed as HR with 95% CI. A sensitivity Cox regression survival analysis was performed for a subset of the unmatched groups separated for N0M0 and N1 or M1 disease, and was also fitted for the matched groups with further adjustment for treatment allocation. Fisher’s exact test and a two-sample z-test of proportions were used to assess differences between treatments received in the matched groups. A secondary analysis assessed biochemical-free and castrate-resistant–free survival within treatment groups. Data for patients who died from PCa or other causes were censored at date of death. All tests were two-sided, with statistical significance set at p < 0.05. All analyses were performed using IBM SPSS for Windows, version 22.0, GraphPad Prism, version 6, and STATA version 14.

3. Results

3.1. ProtecT case and ACN control characteristics

The flow of the patients through the study is summarised in Figure 1. There were 513 ProtecT advanced cases, of whom 21 were excluded because of incomplete data at presentation. For the remaining 492 cases, the mean age was 64 yr (interquartile range [IQR] 61–68); median PSA was 17 ng/ml (mean 33, IQR 8–32 ng/ml); 43% had PSA ≥20 ng/ml; 62% had clinical stage ≥T3; 23% had a Gleason score ≥8; and 8% had N1 or M1 disease. Median follow-up was 7.4 yr (IQR 5.5–9.7; Table 1). For analysis of biochemical recurrence, data on primary treatment were available for 432 out of 492 ProtecT cases (88%), and data on PSA follow-up and on neoadjuvant, adjuvant, or salvage therapies for 352 out of 492 cases (72%).

We identified 3978 ACN controls aged 50–72 yr with clinically detected PCAs. Median follow-up was 5 yr (IQR 3.1–7.8). There were differences in baseline characteristics: ACN controls were older, had higher PSA levels, higher Gleason scores, and higher PCa stages (all p < 0.0002). Accordingly, we matched ProtecT cases (n = 401) to ACN controls (n = 401) across these variables (Table 1). The median follow-up for the matched cohorts was 7.6 yr (IQR 5.1–9.8). There were complete data on primary treatment for 352 of 401 (88%) matched ProtecT cases and 391 of 401 (98%) ACN controls (Table 2).

3.2. Survival analysis

3.2.1. ProtecT advanced cases

Of the 492 ProtecT men, 54 (11%) had radical prostatectomy (RP); 245 (50%) had RT, of whom 93% had neoadjuvant and
adjuvant ADT; 122 (25%) had ADT alone; five (1%) had primary chemotherapy; six (1%) had other treatment (high-intensity focused ultrasound or monitoring); and for 60 (12%) the treatment was unknown. We were unable to demonstrate a difference in PCa-specific (HR 0.95, 95% CI 0.22–4.12; p = 0.94) or all-cause mortality (HR 0.69, 95% CI 0.29–1.67; p = 0.41) between the RP and RT groups (Fig. 2A,2B). Men who received RP were younger (p < 0.01) and had lower PSA (p < 0.0001) compared to the RT group, but no significant difference was observed in Gleason score (p = 0.84) or stage (p = 0.19; Supplementary Table 1).

All-cause mortality was 7% (4/54) among men who underwent RP (2 died of PCa; 4%) and 15% (37/245) among those who received RT (12 died of PCa; 5%). All-cause mortality was 7% (4/54) among men who underwent RP (2 died of PCa; 4%) and 15% (37/245) among those who received RT (12 died of PCa; 5%). All-cause

Table 2 – Primary treatments and death rates among matched ProtecT cases and Anglia Cancer Network (ACN) controls

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Matched ProtecT cases</th>
<th>Matched ACN controls</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N</td>
<td>Deaths, n (%)</td>
</tr>
<tr>
<td></td>
<td>PCS</td>
<td>AC</td>
</tr>
<tr>
<td>RP</td>
<td>47</td>
<td>1 (4)</td>
</tr>
<tr>
<td>RT + ADT</td>
<td>200</td>
<td>11 (6)</td>
</tr>
<tr>
<td>Nonradical</td>
<td>105</td>
<td>19 (18)</td>
</tr>
<tr>
<td>Unknown</td>
<td>49</td>
<td>6 (12)</td>
</tr>
<tr>
<td>Total</td>
<td>401</td>
<td>37 (9)</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>n</td>
<td>Deaths, n (%)</td>
</tr>
<tr>
<td></td>
<td>PCS</td>
<td>AC</td>
</tr>
<tr>
<td>RP</td>
<td>150</td>
<td>5 (3)</td>
</tr>
<tr>
<td>RT + ADT</td>
<td>127</td>
<td>6 (5)</td>
</tr>
<tr>
<td>Nonradical</td>
<td>114</td>
<td>51 (45)</td>
</tr>
<tr>
<td>Unknown</td>
<td>10</td>
<td>1 (10)</td>
</tr>
<tr>
<td>Total</td>
<td>401</td>
<td>63 (16)</td>
</tr>
</tbody>
</table>

RP = radical prostatectomy; ADT = androgen deprivation therapy; PCS = prostate cancer–specific; AC = all causes.

* Adjuvant ADT was given in combination with radical radiotherapy in 93% of ProtecT cases and 88% of ACN controls.

* Nonradical treatment includes primary ADT, palliative chemotherapy, palliative radiotherapy, and monitoring.
mortality was higher among men who underwent non-radical treatment (51/133; 38%) and men whose treatments were unknown (25/60; 42%; all p < 0.0001; Fig. 2A,2B). Men treated using ADT were older (p = 0.01) and had higher PSA (p < 0.0001), Gleason score (p = 0.05), and stage (p < 0.0001) compared to men who received radical treatment (Supplementary Table 1).

3.2.2. Comparison with ACN controls: Kaplan-Meier survival analysis

We found lower risks of death from PCa (HR 0.29, 95% CI 0.38–0.53; p < 0.0001) and from all causes (HR 0.45, 95% CI 0.48–0.63; p < 0.0001) among ProteCT cases compared to unmatched ACN controls (Supplementary Fig. 1A,1B). After matching (Table 1) we observed a 45% lower rate of death from PCa (HR 0.55, 95% CI 0.38–0.83; p = 0.0037), but were unable to demonstrate a difference in all-cause deaths (HR 0.83; 95% CI 0.63–1.1; p = 0.19) between matched ProteCT cases and ACN controls at 7.6 yr (Fig. 3A,3B).

There was a similar proportion of men who received radical and nonradical treatments in the matched groups (p = 0.87), but more men in the ProteCT group received RT compared to the matched ACN controls (p < 0.0001; Table 2).

Among the ProteCT matched cases, 247 men received radical treatment (RP n = 47; RT n = 200) of whom 12 died from PCa (RP n = 1 [4%]; RT n = 11 [6%]) and 33 died of all causes [RP n = 2 [4%]; RT n = 31 [16%]].

Among the ACN matched controls, 277 men received radical treatment (RP n = 150; RT n = 127) of whom 11 died of PCa (RP n = 5 [3%]; RT n = 6 [5%]) and 32 died of all causes (RP n = 12 [8%]; RT n = 20 [16%]).

Among the matched men who received nonradical treatment, a significantly greater proportion died in the ACN control group (n = 114; 51 PCa deaths and 68 all-cause deaths) than in the ProteCT group (n = 105; 19 PCa deaths and 37 all-cause deaths; p < 0.0002; Table 2).

3.2.3. Comparison with ACN controls: Cox proportional hazards survival analysis

Multivariable analysis for the unmatched groups revealed that ProteCT cases (n = 404) had a 53% lower risk of death from PCa (HR 0.47, 95% CI 0.34–0.66; p < 0.0001) and a 30% lower risk of death from all causes (HR 0.70, 95% CI 0.56–0.88; p < 0.0001) compared to unmatched ACN controls (n = 3335; Table 3). Higher PSA, higher Gleason score, and higher stage all indicated a greater risk of death, whereas later years of diagnosis lowered the risk. There was also a higher risk of death from all causes in the oldest age group.

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(67–72 yr). A subset analysis for men with NO, M0 disease did not demonstrate a difference in the risk of death from PCa (HR 0.69, 95% CI 0.45–1.06; p = 0.09) or all causes (HR 0.94, 95% CI 0.73–1.22; p = 0.65) between the unmatched groups. However, men with N1 or M1 disease had a much lower risk of death from PCa (HR 0.33, 95% CI 0.18–0.59; p < 0.0001) and all causes (HR 0.38; 0.22–0.63; P < 0.0001) in the ProtecT group than in the ACN group (Supplementary Table 2).

Multivariable analysis was performed for the matched groups after further adjusting for treatment received. We did not find evidence of a difference in the risk of death from PCa among men who received radical treatment (HR 1.91, 95% CI 0.73–5.02; p = 0.19). Men treated with RT had a higher risk of death from all causes compared to the RP group (HR 2.02, 95% CI 1.08–3.77, p = 0.03). There was a much higher risk of death from PCa (HR 6.70, 95% CI 2.64–16.9; p < 0.0001) and all causes (HR 4.55, 95% CI 2.42–8.52; p < 0.0001) among men who received nonradical treatment compared to those who underwent radical treatment (Supplementary Table 3).

3.2.4. Kaplan-Meier analysis of biochemical recurrence by primary treatment group in the ProtecT group
PSA follow-up was available for ProtecT cases and is reported in more detail in the Supplementary material. At a median of 7.4 yr, PCa-specific survival was 96% in the RP group and 96% in the RT group. There were no predictors of biochemical failure, PCa-specific mortality, or overall mortality among men treated with RP or RT on univariable or multivariable analysis, except for high Gleason score, which increased the risk of death from all causes in the RT group (HR 6.48, 95% CI 1.48–28.4; p = 0.01; Supplementary Table 4 and Supplementary Fig. 2).

4. Discussion
This study reports on asymptomatic men who were excluded from ProtecT because of advanced and high-risk PCa; their outcomes form an important component of the overall context of the ProtecT study and its generalisability with respect to treatment of PSA-detected PCa. In men who were excluded from ProtecT, but were treated radically, we found low rates of all-cause and PCa specific deaths (14% and 5%), with no differences between surgery and radiotherapy at a median of 7.4 years. Most deaths occurred among men receiving nonradical treatments, probably because they had more advanced disease and/or were not fit for radical treatment, although very unfit men were screened out from ProtecT by the general practitioner.

With respect to the main clinical outcome paper from ProtecT, all-cause mortality ((~10% at a median of 10 yr)[13] was lower than that noted here in the RT group (15%). This suggests that ProtecT men with advanced PCa treated by RT in the present study were less fit than those in the randomised group. Moreover, the group who received nonradical treatment and those whose treatment was unknown had significantly greater all-cause mortality (38% and 42%, respectively) compared with those undergoing radical treatment. The PCa-specific mortality among the ProtecT group receiving radical treatment (5%) was greater than that found in the randomised group (~1%), but nevertheless indicates very good cancer survival.

The reduction in risk of death from PCa among advanced ProtecT cases (45%) compared to ACN controls persisted after careful case-control matching to attempt to compensate for leadtime bias and differences in baseline characteristics. However, other biases cannot be ruled out, including the greater number of men undergoing surgery in
the ACN group than in the ProtecT group when comparing those who received radical treatment, and the fact that the ACN group were generally less fit. However, there were no differences in PCa-specific or all-cause mortality between the matched cases and controls when comparing those who received radical treatment. The higher death rates observed among ACN controls occurred mainly in men who received nonradical treatments, suggesting early detection may improve the life expectancy of this subgroup, although other explanations such as group heterogeneity, leadtime bias, and selection bias cannot be ruled out.

Cox regression results for survival analysis (53% lower risk of death from PCa and a 30% reduction in all-cause mortality in the ProtecT group) can probably be explained in part by leadtime bias in the ProtecT cohort [18,19].

We found no difference in PCa-specific or overall survival between the RP and RT ProtecT groups. Only a small proportion of men who received radical treatment (RP 4%, RT 5%) died from PCa, which adds to increasing evidence that radical treatment of locally advanced or high-risk disease delivers good oncologic outcomes[8,9]. The all-cause and PCa survival outcomes for the ProtecT group are better than in most studies on men clinically presenting with advanced disease [20], supporting the hypothesis that early detection of advanced and high-risk PCa may be of benefit. The wider context of the impact of PSA testing on community-based men will be presented in the findings of the CAP (Cluster randomised trial of prostate cancer) trial in 2017 [21].

The quality of data for the ACN group is likely to be good [15,21]. We minimised misattribution of death by using two independent clinicians blinded to the study group and treatment, and by checking with the ProtecT recruitment centre of origin. Matching reduced the number of men for the matched analysis (n = 401) and there may be additional biases that our matching process was unable to take into account. Multidisciplinary teams reviewed the histopathology for ACN controls, whereas ProtecT cases were reviewed by the expert ProtecT histopathology group [22]. For surgically treated ACN cases, however, histology was reviewed centrally. Potential differences in grade and stage allocation may have had some impact on apparent survival benefits among the ProtecT cases. There was no information available on the comorbidity burden for the ACN controls, and therefore we were unable to match the two groups according to these factors. ProtecT participants were 98% Caucasian and patients with a prior history of cancer were excluded, which may have influenced overall survival. The natural history of PCa can be long and further follow-up is required, but such leadtime factors are likely to be of lesser magnitude among men with advanced disease [23–25].

5. Conclusions

PSA testing identifies asymptomatic men with advanced and high-risk PCa whose early treatment leads to good survival rates. We observed improved survival in the ProtecT men who received nonradical treatment compared to men presenting clinically without PSA testing, although leadtime and selection bias are difficult to exclude. It will be important to assess longer-term survival and add patient-reported outcomes among these men to assess the balance between treatment impact and survival.

Author contributions: Alastair D. Lamb had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

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Analysis and interpretation of data: Johnston, Parashar, Shaw, Lamb, Xiong.

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Statistical analysis: Johnston, Parashar.

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Appendix A. Supplementary data

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