BMJ Open Diagnostic and prognostic factors in patients with prostate cancer: a systematic review

Katharina Beyer ^(b), ¹ Lisa Moris, ² Michael Lardas, ³ Anna Haire, ¹ Francesco Barletta ^(b), ⁴ Simone Scuderi, ⁴ Megan Molnar ^(b), ⁵ Ronald Herrera, ⁵ Abdul Rauf, ⁶ Riccardo Campi ^(b), ⁷ Isabella Greco, ⁷ Kirill Shiranov, ⁸ Saeed Dabestani, ⁹ Thomas van den Broeck, ² Sujenthiran Arun, ¹⁰ Mauro Gacci, ⁷ Giorgio Gandaglia, ⁴ Muhammad Imran Omar ^(b), ¹¹ Steven MacLennan ^(b), ¹¹ Monique J Roobol, ¹² Bahman Farahmand, ¹³ Eleni Vradi, ⁵ Zsuzsanna Devecseri, ¹⁴ Alex Asiimwe, ⁵ Jihong Zong, ¹⁵ Sara J Maclennan, ¹¹ Laurence Collette, ¹⁶ James NDow, ¹² Alberto Briganti, ^{4,17} Anders Bjartell, ¹⁸ Mieke Van Hemelrijck, ¹ and the PIONEER Consortium

ABSTRACT

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For numbered affiliations see end of article.

Correspondence to Katharina Beyer;

Katharina Beyer; katharina.beyer@kcl.ac.uk **Objectives** As part of the PIONEER Consortium objectives, we have explored which diagnostic and prognostic factors (DPFs) are available in relation to our previously defined clinician and patient-reported outcomes for prostate cancer (PCa).

Design We performed a systematic review to identify validated and non-validated studies.

Data sources MEDLINE, Embase and the Cochrane Library were searched on 21 January 2020.

Eligibility criteria Only quantitative studies were included. Single studies with fewer than 50 participants, published before 2014 and looking at outcomes which are not prioritised in the PIONEER core outcome set were excluded.

Data extraction and synthesis After initial screening, we extracted data following the Checklist for Critical Appraisal and Data Extraction for Systematic Reviews of prognostic factor studies (CHARMS-PF) criteria and discussed the identified factors with a multidisciplinary expert group. The quality of the included papers was scored for applicability and risk of bias using validated tools such as PROBAST, Quality in Prognostic Studies and Quality Assessment of Diagnostic Accuracy Studies 2.

Results The search identified 6604 studies, from which 489 DPFs were included. Sixty-four of those were internally or externally validated. However, only three studies on diagnostic and seven studies on prognostic factors had a low risk of bias and a low risk concerning applicability.

Conclusion Most of the DPFs identified require additional evaluation and validation in properly designed studies before they can be recommended for use in clinical practice. The PIONEER online search tool for DPFs for PCa will enable researchers to understand the quality of the current research and help them design future studies.

Ethics and dissemination There are no ethical implications.

Strengths and limitations of this study

- A multidisciplinary team including patients, urologists, oncologists, radiation oncologists, methodological experts and pathologists were involved throughout the study.
- The search was restricted from 2014 onwards, to maintain a pragmatic approach.
- The main strength of this study is the extensive and comprehensive search and screening of the studies included.

INTRODUCTION

Prostate cancer (PCa) accounts for 15% of cancers diagnosed¹ and is the second most common cancer in males worldwide.² PCa is clinically and molecularly heterogeneous and is usually suspected based on the clinical findings of digital rectal examination and/ or prostate-specific antigen (PSA) levels.¹ However, which diagnostic or prognostic factors (DPFs) can be used to select patients for specific therapeutic options remains largely unclear.³ Specific biomarkers in urine or in blood are available on top of traditional PSA testing, such as PCA3, TMPRSS2-ERG fusion or kallikreins as incorporated in the Phi or 4Kscore test together with other parameters including family history.4-7 However, the European Association of Urology (EAU) guidelines (2021) currently do not provide general recommendations to implement these biomarkers into routine screening programmes due to limited data. As part of the American Society of Clinical Oncology (ASCO) guidelines, Eggener et al recommended commercially available biomarkers, which have been shown to provide prognostic significance and additional information beyond standard clinical models in patient selection in the localised context: Oncotype Dx Prostate, Prolaris, Decipher, and ProMark.⁸ However, no guidelines have recommended DPFs for other stages of PCa. The expert panel at the Advanced Prostate Cancer Consensus Conference (APCCC) consensus meeting of advanced PCa in Basel 2019, recommended AR-V7 for mCRPC as potentially useful, which ultimately led to the inclusion of AR-V7 testing in the NCCN guidelines.⁹

The PIONEER Consortium is an international collaboration coordinated by the EAU, which aims to establish the best evidence-based management and clinical practice of PCa across all disease stages using the power of big data analytics towards a more outcome-driven, value-based and patient-centric healthcare system.¹⁰ A key objective is to address one of the major challenges within the context of diagnostic or prognostic biomarkers/factors: the inability to incorporate DPFs into the management of PCa in terms of screening, diagnosis and treatment. It is therefore important to summarise and evaluate the evidence. Biomarkers can be classified into different types: diagnostic, prognostic, predictive and therapeutic-in this study we focus on the first two.¹¹ A diagnostic biomarker or factor is useful when cancer is suspected and allows the early detection based on symptoms or tests.¹¹ The overall aim of a diagnostic biomarker is to distinguish people with the diseases from people without the disease. A prognostic biomarker or factor is a clinical or biological characteristic which provides information on the likely course of the disease, that is, biochemical progression or disease recurrence.¹¹ It enables clinicians to decide on the most suitable treatment depending on the likely course of the disease. In the sections below, we have used the terms biomarkers and factors interchangeably. Multiple DPFs can be measured in tissue, blood or urine. These come with different advantages and disadvantages and only a limited number of factors are currently available for PCa in standard clinical care.

We aimed to systematically review the evidence from 2014 onward to assess which DPFs are available in relation to previously defined outcomes for PCa.

METHODS

The systematic review (SR) followed the Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines.¹² A detailed protocol of the overall project was published elsewhere¹³ (please see the protocol attached as methods online supplemental appendix). Briefly, we followed the following four steps (figure 1):

1. Comprehensive systematic literature review of DPFs for all stages of PCa (localised, locally advanced, metastatic, and non-metastatic castration resistant) from 2014 onwards. DPFs developed before 2014 were not included, due to the significant changes influencing the staging of PCa (i.e., Consensus Conference on

| Workflow | Task |
|----------|--|
| Stage 1. | Broad literature-based systematic review of diagnostic and prognostic factors (DPs) for all stages of prostate cancer from 2014 onwards (English only; humans). • Extract data from the included studies following the CHARMS-PF guideline. |
| Stage 2. | Discussion of systematic review findings by a multidisciplinary expert panel Review the list of included studies |
| Stage 3. | Risk of Bias Assessment and applicability of individual studies using PROBAST, QUIPS and QUADAS-2 |
| Stage 4. | Quantitative assessment of individual articles using meta-analytic techniques: If PROBAST indicates low risk of bias and low concerns for applicability: Oxford Classification Centre for Evidence Based Medicine: If there is Level 1a (SR of RCTs), we do not do a meta-analysis No Level 1a but >2 RCTs, we do a meta-analysis No Level 1a/b, i.e. if at least two RCTs are now available, and systematic review of RCT evidence is not possible, we will identify whether there is a systematic review for observational studies (real world evidence; RWE), we do not do a meta-analysis If systematic review of RWE is not available, a systematic review of observational study will be conducted, and a meta-analysis will be performed if at least two RWE studies are available and data pooling is feasible and there are low concerns of risk of bias. |

Final aim: Develop online PIONEER Online Search Tool for DPFs

Figure 1 Overview of four stage process. CHARMS-PF, Checklist for Critical Appraisal and Data Extraction for Systematic Reviews of prognostic factor studies; DPFs, diagnostic and prognostic factors; PROBAST, Prediction model Risk Of Bias Assessment Tool; QUADAS-2, Quality Assessment of Diagnostic Accuracy Studies 2; QUIPS, Quality in Prognostic Studies; SR, systematic review.

Gleason Grading of Prostatic Carcinoma [60]) that have taken place in diagnostic and prognostic practice and patient management since then.

- 2. Assessment and identification of final list of DPFs by a multidisciplinary expert panel.
- Evaluation of quality of studies published using risk of bias (RoB) tools: Prediction model RoB Assessment Tool (PROBAST) if applicable; or Quality in Prognostic Studies (QUIPS) tool for prognostic and the Quality Assessment of Diagnostic Accuracy Studies 2 (QUA-DAS-2) tool for diagnostic factors.
- 4. Due to the heterogeneity of the studies identified no further formal quantitative assessments in the form of a meta-analyses could be performed. Hence, the findings of stages 1–3 have been reported here as the results of a SR.

Stage 1: comprehensive literature review

We developed the search criteria for the first search with an information scientist who specialises in SR for urology. MEDLINE, Embase and the Cochrane Library were searched on 21 January 2020. The second search was developed following a consultation with an independent information scientist group who excluded row 12, 14 and 16 of (see online supplemental table 1). We screened the EAU Guidelines reference list for PCa in our third search (see figure 2).

Stage 2: multidisciplinary expert meeting

On the 20 March 2020, we invited a group of multidisciplinary participants to discuss the identified articles on DPFs (see online supplemental table 2). The participants were presented the search criteria and the extracted data. Data extraction followed the CHARMS-PF checklist and we added author and year of publication.

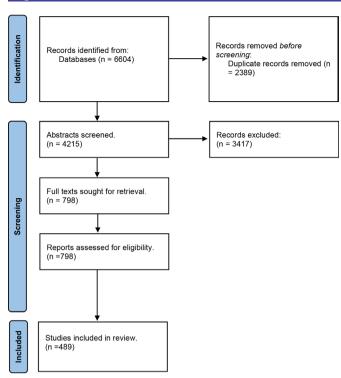


Figure 2 PRISMA flow diagram. PRISMA, Preferred Reporting Items for Systematic Reviews and Meta-Analyses; COS, Core outcome set.

Stage 3: evaluation of quality of studies published using the RoB tools

Prior to the evaluation of the quality of studies, an initial pilot screening to prepare the raters for the use of PROBAST, QUADAS-2 and QUIPS was performed. This aimed to reach consensus on how to judge the domains of the assessments using the three RoB tools. Two urologists (FB and SS) and two epidemiologists (AH and KB) were involved in the pilot assessments. The group discussed any discrepancies. Articles which presented the development and validation either internal validation or external validation (i.e., the same data was used for both development and internal validation, such as bootstrapping or cross-validation; different populations were used for development and validation), of a diagnostic or prognostic model were assessed with PROBAST. Papers assessing single biomarkers or with/without validation were assessed with QUIPs for prognostic or QUADAS-2 for diagnostic biomarkers.

Evaluation of quality of studies published using QUADAS-2

The RoB of diagnostic factors without validation or single validated factors was evaluated using QUADAS-2. We assessed the following four domains: patient selection, index test, reference standards and flow and timing. The first three domains are assessed looking at applicability and all four domains were assessed in terms of RoB.¹⁴ We created a summative score after the diagnostic studies were assessed by two reviewers and in case of disagreement a third reviewer assessed the study.

Evaluation of quality of studies published using PROBAST (diagnostic)

The RoB of internal or external validated diagnostic models was assessed using the PROBAST RoB tool. PROBAST includes four domains assessing the RoB (i.e., participants, predictors, outcome, and analysis) and four domains assessing applicability (i.e., participants, predictors, and outcome) (see online supplemental table 3 for scoring information).

Evaluation of quality of studies published using QUIPS

To assess the articles which are single factors or were not internally or externally validated, we used the QUIPS rating procedure (see online supplemental table 4 for scoring information). To standardise the approach across raters, we used the QUIPS electronic spreadsheet (excel) from Hayden et al.¹⁵ There are no rules available for QUIPS on how to score the overall RoB of a paper. Due to the large number of papers and the need for synthesis, we followed the suggestions from Grooten et al., and categorised on the following criteria: (1) Paper was classified as low RoB if all domains were classified as having low RoB, or up to one moderate RoB; (2) Paper was classified as high RoB if one or more domains were classified as having high RoB, or \geq 3 moderate RoB; (3) Paper was classified as having moderate RoB if all papers in between 1 or 2 (see online supplemental table 1). This assessment was based on the risk scores of individual assessments within the group. If the overall assessment was not possible due to differences in the individual category, a third assessor reviewed the assessments and the results were discussed.

Evaluation of quality of studies published using PROBAST (prognostic)

The RoB of prognostic validated models were assessed using PROBAST. As highlighted above, PROBAST includes four domains assessing the RoB (i.e., participants, predictors, outcome and analysis) and the domains assessing applicability (i.e., participants, predictors and outcome).

RESULTS

Stage 1: comprehensive literature review

Stage 1 identified 6604 citations and contained three independent searches. After removing duplicates, we screened 4215 abstracts, from which 489 met the inclusion criteria.

Stage 2: multidisciplinary expert meeting

The group discussed the results and additional literature on DPFs was suggested to help the classification of the DPFs, such as the ASCO Guideline on Molecular Biomarkers in Localised Prostate Cancer.¹⁶

Stage 3: evaluation of quality of studies published using the RoB tools

The 489 articles were equally divided between six groups. The six groups received the guidance documents which

| Table 1 Overall judgement of RoB QUADAS-2, diagnostic Image: Control of | | | | | | |
|--|-----|---------------|--|--|--|--|
| Overall judgement of RoB | RoB | Applicability | | | | |
| Low | 10 | 10 | | | | |
| High | 23 | 21 | | | | |
| Unclear | 8 | 10 | | | | |
| Total | 41 | | | | | |
| PROBAST, diagnostic | | | | | | |
| Overall judgement of RoB | RoB | Applicability | | | | |
| Low | 3 | 8 | | | | |
| High | 14 | 10 | | | | |
| Unclear | 3 | 2 | | | | |
| Total | 20 | | | | | |
| QUIPS | | | | | | |
| Overall judgement of RoB | RoB | | | | | |
| Low | 29 | | | | | |
| Moderate | 49 | | | | | |
| High | 307 | | | | | |
| Total | 385 | | | | | |
| PROBAST, prognostic | | | | | | |
| Overall judgement of RoB | RoB | Applicability | | | | |
| Low | 3 | 15 | | | | |
| High | 27 | 20 | | | | |
| Unclear | 13 | 8 | | | | |
| Total | 43 | | | | | |

PROBAST, Prediction model Risk Of Bias Assessment Tool; QUADAS-2, Quality Assessment of Diagnostic Accuracy Studies 2; QUIPS, Quality in Prognostic Studies; RoB, risk of bias.

were identified during the pilot phase.^{14 15 17-19} In addition, MvH and KB discussed questions with each individual group.

Evaluation of quality of studies published using QUADAS-2

The RoB of the 41 included studies was low for 10 studies, high for 23 studies and unclear for eight. RoB concerning applicability was low for 10 studies, high for 21 studies and unclear for 10 studies (see table 1). Table 2 shows the studies with an overall low RoB across both categories. Two studies were identified to have an overall low RoB.^{20 21}

Evaluation of quality of studies published using PROBAST (diagnostic)

We identified 20 papers to be assessed with PROBAST. The RoB of three papers was low, high for 14 and was unclear for three. The applicability of eight papers was high and was unclear for two (see table 1). Online supplemental table 1 shows the criteria on how to judge the RoB. One study had an overall low RoB across both domains. All categories except 'predictors' was scored to have a low RoB. There was little information available for the category predictors and therefore it was scored as 'unclear' (see table 3).

Evaluation of quality of studies published using QUIPS

The 12 assessors independently inserted the relevant information and assessed each domain such as participation, attrition, prognostic factor confounding and statistical analysis and reporting.

A total of 387 prognostic factors were assessed using QUIPs. A total of 307 papers were classified as high RoB. Forty-nine papers were classified as having a moderate RoB and 28 papers were scored as low RoB (see table 1). Out of the 28 papers with a low RoB, the most common moderate bias was linked to attrition (12 papers), followed by confounding (4 papers), participation (3 papers), outcome (1 paper), statistical analysis (1 paper) (see table 4).

Evaluation of quality of studies published using PROBAST (prognostic)

The assessors identified 44 papers to be assessed with PROBAST, of those three scored a low RoB, 27 a high RoB and 13 were assessed as unclear (see table 1). In terms of applicability, 15 papers scored low, 20 high and eight unclear. Two papers were scored to have an overall low $RoB^{22\,23}$ (see table 3).

Characteristics of studies identified with low RoB

Details of the identified validated DPF models with an adequate quality are presented in table 5. We identified 32 studies with an overall low RoB (assessed with PROBAST, QUIPS, QUADAS-2). Out of these 32 studies, we identified one validated diagnostic model (assessed with PROBAST),²⁴ two validated prognostic models (assessed with PROBAST),^{22 23} two non-validated diagnostic single factors (assessed with QUADAS-2)^{20 21} and 26 prognostic factors (assessed with QUIPS)²⁰⁻⁵⁰ which have not been validated and two single prognostic factors which have been validated (assessed with QUIPS).^{34 50} Prognostic factors assessed with QUIPS were identified with a low RoB for the localised PCa population. Sixtyseven per cent of the low RoB DPFs were intended to be measured after the treatment was performed. In addition,

| Table 2 Nor | Table 2 Non-validated DPFs with overall low RoB: QUADAS-2 | | | | | | | | | |
|------------------------|---|-------------------|------------------|--------------------|-----------------|-------------------|------------------|--------------------|-----|---------------|
| Author | Year | Patient selection | Index test(s) | Reference standard | Flow and timing | Patient selection | Index test(s) | Reference standard | RoB | Applicability |
| Hagiwara ²⁰ | 2017 | Low | Low | Low | Low | Low | Low | Low | Low | Low |
| Kelly ²¹ | 2015 | Low | Low | Low | Low | Low | Low | Low | Low | Low |

DPFs, diagnostic and prognostic factors; QUADAS-2, Quality Assessment of Diagnostic Accuracy Studies 2; RoB, risk Of bias.

| | ROB | | | | Applicability | | | Overall | |
|--------------------------|--------------|------------|---------|----------|---------------|------------|---------|---------|---------------|
| Author | Participants | Predictors | Outcome | Analysis | Participants | Predictors | Outcome | ROB | Applicability |
| Diagnostic | | | | | | | | | |
| Guinney ²² | Low | Low | Low | Low | Low | Low | Low | Low | Low |
| Joniau ²³ | Low | Low | Low | Low | Low | Low | Low | Low | Low |
| Prognostic | | | | | | | | | |
| Palsdottir ²⁴ | Low | Unclear | Low | Low | Low | Low | Low | Low | Low |

DPFs, diagnostic and prognostic factors; PROBAST, Prediction model Risk Of Bias Assessment Tool; RoB, risk of bias.

the most commonly measured outcome was biochemical recurrence followed by overall survival. However, it is important to take into consideration that even from the studies assessed with a low RoB, only 2 out of the 32 were of a non-observational study design.

As highlighted above, we identified three validated DPFs which were scored to have a low RoB and low risk concerning applicability. First, we identified the 'Unified Prostate Cancer Risk Prediction Model Combining the Stockholm 3 Test and MRI', a risk prediction model which combines clinical variables, genetic and protein biomarkers. Five hundred and thirty-two men were involved across three centres.²⁴ Second, the DREAM challenge developed a set of five standardised raw event-level tables, using laboratory values, patients' demographic information, medical history, lesion sites, previous treatments and vital signs of patients with mCRPC. These variables where combined by using data from four clinical trials.²² Third, Joniau *et al.*, developed 'Pretreatment Tables' to predict the pathologic stage of locally advanced PCa after RP based on pretreatment PSA level and biopsy Gleason score.²³

We identified two single factors which were validated and had low RoB. First, Lara *et al.*, assessed and validated the serum biomarkers of bone metabolism (N-telopeptide and pyridinoline) and formation (C-terminal collagen propeptide and bone alkaline phosphatase)) in 778 CRPC patients as part of the randomised phase III SWOG trial (S0421) of docetaxel/prednisone with or without atrasentan.³⁴ Second, Berg *et al.*, showed that ERG expression can be used to estimate the risk of progression during AS including 265 patients at diagnosis and progression during AS.⁵⁰

DISCUSSION

Despite the large number of studies on DPFs which are published every year, there is a paucity of DPFs that are suitable to be incorporated into clinical practice. The majority of DPFs have not yet been validated and are identified in poor quality studies. Our analysis found that most identified studies had a high to moderate RoB due to poor design standards, conduct, reporting and/or analysis that is, generalisability and size of the population, poor model development (no testing or missing important confounders) or only correlation studies, missing data was rarely reported. However, we did identify a small number of validated DPFs with low RoB. We identified three validated models which combine: first, clinical variables, genetic and protein biomarkers, and improved clinical outcome performance of PCa diagnostics (The Unified Prostate Cancer Risk Prediction Model)²⁴; second, laboratory values, patients' demographic information, medical history, lesion sites, previous treatments and vital signs of patients with metastatic castration-resistant PCa (DREAM challenge)²²; and third, pretreatment PSA level and biopsy Gleason score to predict the pathological stage of locally advanced PCa ('Pretreatment Tables').²³

Two single factors have been validated: the serum biomarkers of bone metabolism in CRPC patients³⁴ and the ERG expression, which can be used to estimate the risk of progression during AS,⁵⁰ which has already been highlighted in the clinical guidelines.¹

Aladawani *et al.*, assessed prediction models for PCa to be used in primary care settings in their SR and identified five models which met their inclusion criteria. From these identified models only one model was externally validated and only one (the Lazzari model 2^{51} had the potential to be implemented in primary care. Lazzari *et al.*, had the lowest RoB (based on PROBAST); however, it must be externally validated before it can be implemented. Hence, Aladawani *et al.*, also concluded that the existing models have limitations concerning study design and reporting performance.⁵²

Tian *et al.*, conducted a review on biomarkers for CRPC patients, however, their quality assessment was focused on study design (RCT vs. observational study), whereas we focused on biomarker specific tools.⁵³ While Tian *et al.*, and our review identified similar factors and quality scores, there were slight discrepancies between the overall RoB assessments. Tian *et al.*, used an overall quality assessment scale from 1 to 6 instead of low, medium and high. In their assessment the validated prognostic study by Lara *et al.*,³⁴ and the non-validated prognostic factor by Pei *et al.*,⁴¹ were scored on the quality scale as 4 (medium quality). We assessed Lara *et al.*,³⁴ to have a low RoB with a moderate risk of confounding and Pei *et al.*,⁴¹ with a moderate RoB concerning the prognostic factor itself. This might explain the discrepancies between the two

| Author | Year | RoB | Population | Study design | Timing | Index | Outcomes |
|-------------------------|------|--------------------|-------------------------|---------------------|--------------------------|---|---|
| alsdottir ²⁴ | 2019 | Diag. | Localised PCa | Observational study | | S3M-MRI (Stockholm3 +PIRADS) | csPCa diagnosis |
| | 2010 | PROBAST | | | The treatment | · · · · | , C |
| iuinney ²² | 2017 | Prog. PROBAST | mCRPC | RCT | Post treatment | ePCR model | OS |
| oniau ²³ | 2017 | Prog. PROBAST | Locally advanced PCa | Observational study | Post-treatment | Gleason score +PSA | Adverse pathologica features at RP; LNI |
| lagiwara ²⁰ | 2017 | QUADAS | Localised PCa | Observational study | Pre-treatment | WFA-reactive glycan-carrying PSA- Gi | PCa diagnosis, PSA- free survival |
| Kelly ²¹ | 2015 | QUADAS | Localised PCa | Observational study | Pre-treatment | miR-141, -145, -155, let7a | PCa diagnosis |
| guilera ²⁵ | 2015 | QUIPS | High risk PCa | Observational study | Pre and post treatment | Age, rectal examination, PSA, biopsy Gleason score, uni/ bilateral tumour, affected cylinder percentage) and postoperative | BCR |
| Alvim ²⁷ | 2019 | QUIPS | Metastatic PCa | Observational study | Post-treatment | PSA response (PSA reduction ≥50%) | OS, PFS |
| Bramhecha ²⁶ | 2019 | QUIPS | Localised PCa | Observational study | Post-treatment | PTEN deletion | BCR |
| Bruce ²⁸ | 2016 | QUIPS | Localised PCa | Observational study | Post-treatment | AZGP1 expression | BR-free survival, CR-free survival, PC-specific death |
| rancini ²⁹ | 2018 | QUIPS | mHSPC | Observational study | Post-treatment | Volume | OS, time to CRPC |
| lamada ³⁰ | 2016 | QUIPS | High risk PCa | Observational study | Post-treatment | PSA, PSA density (PSAD), PSAD of the transition zone, percentage of positive cores (PPC), prostate volume, TZ volume, Gleason score, PPC from the dominant side | BCR |
| lashimoto ³¹ | 2020 | QUIPS | Localised PCa | Observational study | Post-treatment | Micro-lymphatic invasion, Gleason | BCR |
| lung ⁵⁷ | 2017 | QUIPS | mCRPC | Observational study | Post-treatment | Neurovascular bundle preservation, blood loss, pT stage, pN stage, pGS, PNI, angiolymphatic invasion, tumour amount in specimen, ECE, PSM, SVI, Bladder neck invasion, Foley duration, post-op undetectable PSA | BCR |
| Kato ³² | 2018 | QUIPS | High risk PCa | Observational study | Post-treatment | LC/IDC | PFS, CSS |
| luth ³³ | 2014 | QUIPS | Localised PCa | Observational study | Post-treatment | No of lymph nodes | BCR |
| ara ³⁴ | 2014 | QUIPS Validated | mCRPC | RCT | Post treatment | Bone resorption and formation | OS |
| .ee ³⁵ | 2016 | QUIPS | Localised PCa | Observational study | Post treatment | Positive surgical margin status and bilateral seminal vesicle invasion | BCR |
| évesque ³⁶ | 2019 | QUIPS | Localised PCa | Observational study | Post treatment | UGT2B17 expression | BCR |
| .in ³⁷ | 2017 | QUIPS | Localised PCa | Observational study | Post treatment | Aberrant Promoter Methylation of Protocadherin8 (PCDH8) | BCR-free survival |
| öffeler ³⁸ | 2015 | QUIPS | mCRPC | Observational study | Anytime | PSA doubling time, PSA nadir during ADT, haemoglobin and alkaline phosphatase levels at CRPC | OS |
| Varang ³⁹ | 2017 | QUIPS | Localised PCa | Observational study | Anytime | PSA: End-of-radiation PSA | BCR-free survival, MFS, CSS, OS |
|)zden ⁴⁰ | 2017 | QUIPS | Localised PCa | Observational study | Post treatment | Age | RRP specimen, BCR, and BCR-free survival rates |
| Pei ⁴¹ | 2016 | QUIPS | CRPC | Observational study | Pre and during treatment | Neutrophil-to-lymphocyte ratio | OS, PFS |
| λu ⁴² | 2016 | QUIPS | mPCa and CRPC | Observational study | Pre treatment | AR-V7 | Time to CRPC / CRPC: CSS |
| ี่ วัน ⁴³ | 2017 | QUIPS | PCa | Observational study | Pre and during treatment | AR-V7 | OS |
| Rüenauver ⁴⁴ | 2014 | QUIPS | Localised PCa | Observational study | Post treatment | YWHAZ | OS |

| Author | Year | RoB | Population | Study design | Timing | Index | Outcomes |
|--------------------------|------|--------------------|------------------------------|---------------------|----------------|--|--|
| Shimodaira ⁴⁵ | 2020 | QUIPS | Metastatic PCa | Observational study | Post treatment | Value of platelet counts | Disease specific survival |
| Strand ⁴⁶ | 2015 | QUIPS | Localised PCa | Observational study | Post treatment | 5-hydroxymethylcytosine score | BCR |
| Takagi ⁴⁷ | 2017 | QUIPS | Localised PCa | Observational study | Post treatment | Age, T stage, % of pos cores, Gleason score, PSA, Total ADT | BCR-free survival |
| Wang ⁴⁸ | 2016 | QUIPS | PCa | Observational study | Post treatment | Platelet to lymphocyte ratio (PLR) | PLR with PFS, CSS and OS n/a |
| Zacho ⁴⁹ | 2017 | QUIPS | Localised PCa | Observational study | Anytime | Bone scan index | Time to CRPC |
| Berg ⁵⁰ | 2014 | QUIPS validated | Under Active Surveillance | Observational study | | ERG immunohisto-chemical staining | Overall AS progression, histopathologic progression |

ADT, androgen deprivation therapy; AS, Active Surveillance; BCR, biochemical recurrence; CSS, cancer-specific survival; DFPs, diagnostic and prognostic factors; mCRPC, metastatic castration resistant prostate cancer; n/a, not available; OS, overall survival; PCa, prostate cancer; PFS, progression-free survival; PI-RADS, Prostate Imaging Reporting and Data System; PROBAST, Prediction model Risk Of Bias Assessment Tool; PSA, Prostate Specific Antigen; PTEN, Phosphatase and tensin homolog; QUADAS, Quality Assessment of Diagnostic Accuracy Studies; QUIPS, Quality in Prognostic Studies; RCT, Randomised control trial; RoB, risk of bias; WFA, Wisteria floribunda agglutinin.

quality assessments. The reports by Alvim *et al.*, Qu *et al.*, were assessed to have the highest quality by Tian *et al.*, ⁵³ similar to our review. This illustrates that different quality assessment tools emphasise different criteria, which may result in small discrepancies. However, the overall conclusion for prognostic single factors was similar in our review and to the work of Tian *et al.*, ⁵³

Similar issues have been identified for other urological cancers. For example, in kidney cancer, a large body of research was identified by Harrison *et al.*, with very few validated studies and lots of heterogeneity.⁵⁴ Schmitz-Dräger *et al.*, published an International Consultation of Urologic Disease/WHO Consensus manuscript where they identified that in bladder cancer one of the main limitations for the lack of incorporation of modern bladder cancer tests into clinical practice decision making is linked to the scarcity of 'good clinical practice guidelines' for the evaluation of diagnostic markers.

There is a need for improved guidance on development and validation of diagnostic markers.⁵⁵ To meet that need, we are developing the PIONEER DPF search tool, which will help researchers and clinicians to get a better understanding of the DPFs for PCa. The tool will not only summarise all relevant studies, but also provide information on the use and results of different RoB assessment tools, which will enable an understanding of the quality of published studies.

Future research should, therefore, focus on addressing the identified shortcomings such as heterogeneity, validation and poor RoB by designing more robust studies which consistently include RoB assessments such as PROBAST, QUIPS or QUADAS-2.

With the growing number of various therapeutic options, diagnosis and management of PCa requires an individualised approach to patient care. There is an unmet need for DPFs to guide decisions for optimal treatment and to predict which patients will benefit the most, from a particular management strategy. DPFs could

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potentially enhance the quality of patient counselling, but currently most need additional evaluation and validation in properly designed studies. Our SR highlights the need for well-designed Real-World Evidence studies, while the PIONEER online search tool can inform the design of new research studies, through providing a rigorous evaluation of the methodological quality of the studies.

The main strength of this study are the extensive and comprehensive search and screening of the studies included. In addition, we are developing an online search tool which showcases the identified and assessed studies. It provides an overview of the available DPFs and enables interested stakeholders to search for DPFs. To our knowledge, this is the first study which has been performed with this extensive amount of literature.

Patient and public involvement

This project has been overseen by a multistakeholder group part of the PIONEER Consortium. PIONEER brings together 35 key stakeholders from academic institutions, patient advocacy groups, European organisations, experts in legal data management, clinicians and pharmaceutical companies, as well as regulatory agencies, economics and ethics, and information and technology specialists. Patients and their family members are therefore involved and actively participate as an integral part of all research conducted by the PIONEER Consortium.

Limitations

Even though this review included three searches and assessments by a multidisciplinary group of fourteen researchers, we recognise potential limitations. Studies were only included from 2014 onwards and DPFs developed before 2014 were not included. However, significant changes which influence the staging of PCa (i.e., Consensus Conference on Gleason Grading of Prostatic Carcinoma⁵⁶ have taken place in diagnostic and prognostic practice and patient management. This changed

Table 5 DPFs

Study Aguilera²⁵ Alvim²⁷ Bramhecha²⁶ Bruce²⁸ Francini²⁹ Hamada³⁰ Hashimoto³ Hung⁵⁷ Kato³² Kluth³³ Lara³⁴ Lee³⁵ Levesque³⁶ Lin³⁷ Loffeler³⁸ Narang³⁹ Ozden⁴⁰ Pei⁴¹ Qu⁴² Qu⁴³ Rizzardi⁵⁸ Ruenauver⁴⁴ Shimodaira⁴ Strand⁴⁶ Takagi47 Wang⁴⁸ Zacho⁴⁹ Berg⁵⁰

| with lov | v risk of bias ass | sessed with | QUIPS | | | | |
|----------|--------------------|-------------|----------------------|------------|-------------|------------------------------------|---------------|
| | Biases | | | Applicabil | ity | | |
| Time | Participation | Attrition | Prognostic factor | Outcome | Confounding | Statistical analysis and reporting | Overall score |
| 2015 | Low | Low | Low | Low | Moderate | Low | Low |
| 2019 | Low | Low | Low | Low | Low | Low | Low |
| 2019 | Low | Moderate | Low | Low | Low | Low | Low |
| 2016 | Low | Moderate | Low | Low | Low | Low | Low |
| 2018 | Low | Low | Low | Low | Low | Moderate | Low |
| 2016 | Low | Low | Low | Moderate | Low | Low | Low |
| 2020 | Low | Low | Low | Low | Low | Low | Low |
| 2017 | Moderate | Low | Low | Low | Low | Low | Low |
| 2018 | Low | Moderate | Low | Low | Low | Low | Low |
| 2014 | Low | Moderate | Low | Low | Low | Low | Low |
| 2014 | Low | Low | Low | Low | Moderate | Low | Low |
| 2016 | Low | Moderate | Low | Low | Low | Low | Low |
| 2019 | Low | Moderate | Low | Low | Low | Low | Low |
| 2017 | Low | Moderate | Low | Low | Low | Low | Low |
| 2015 | Low | Low | Low | Low | Low | Low | Low |
| 2017 | Low | Moderate | Low | Low | Low | Low | Low |
| 2017 | Moderate | Low | Low | Low | Low | Low | Low |
| 2016 | Low | Low | Moderate | Low | Low | Low | Low |
| 2016 | Low | Low | Low | Low | Low | Low | Low |
| 2017 | Low | Low | Low | Low | Low | Low | Low |
| 2015 | Low | Low | Low | Low | Low | Low | Low |
| 2014 | Low | Moderate | Moderate | Low | Low | Low | Low |
| 2020 | Low | Moderate | Low | Low | Low | Low | Low |
| 2015 | Low | Moderate | Low | Low | Low | Low | Low |
| 2017 | Low | Low | Low | Low | Moderate | Low | Low |
| 2016 | Low | Moderate | Low | Low | Low | Low | Low |
| 2017 | Moderate | Low | Low | Low | Moderate | Low | Low |
| 2014 | Low | Low | Low | Low | Low | Low | Low |

DPFs, diagnostic a

the staging of the patient population and therefore has an impact on DPFs.

In addition, there is a potential of subjectivity in the evaluation of the studies. Even though the studies have been assessed in duplicate, there might be variation across groups. However, given the overall moderate to high RoB, this does not influence the overall recommendation of the project.

CONCLUSION

At present DPFs that are capable of significantly improving diagnosis and prognosis in PCa are an unmet need as most of the DPFs identified require additional evaluation and validation in properly designed studies before they can be recommended for use in clinical practice.

Well-designed real world evidence (RWE) studies can help to increase quality. Our SR aims to inform clinicians and patients about this rapidly evolving field, while the PIONEER online search tool for DPFs for PCa will enable researchers to perform future research, and to understand the quality of the current available studies.

Author affiliations

¹Translational and Oncology Research (TOUR), King's College London, Faculty of Life Sciences and Medicine, London, UK

- ²Department of Urology, University Hospitals Leuven, Leuven, Belgium
- ³Department of Urology, Metropolitan Hospital, Athens, Greece

⁶Department of Urology, Mid Cheshire Hospitals, NHS Foundation Trust, Crewe, UK ⁷Department of Minimally Invasive and Robotic Urologic Surgery and Kidney Transplantation, University of Florence, Florence, Italy

⁴Unit of Urology/Division of Oncology, URI, IRCCS Ospedale San Raffaele, Milan, Italy ⁵Bayer AG, Berlin, Germany

9

⁸CDC Zdorovie, Rostov-on-Don, Russia

⁹Dept. of Translational Medicine, Division of Urological Cancers, Lund University, Kristianstad Central Hospital, Malmo, Sweden

¹⁰Flatiron Health, London, UK

- ¹¹Academic Urology Unit, University of Aberdeen, Aberdeen, UK
- ¹²Department of Urology, Erasmus University Medical Center, Rotterdam, The Netherlands
- ¹³Global Epidemiology, Bayer AG, Stockholm, Sweden
- ¹⁴Sanofi, Paris, France

¹⁵Global Medical Affairs Oncology, Real World Evidence, Bayer HealthCare Pharmaceuticals Inc, Whippany, New Jersey, USA

- ¹⁶EORTC Headquarters, Brussels, Belgium
- ¹⁷Department of Urology, University Vita e Salute-San Raffaele, Milan, Italy
- ¹⁸Department of Translational Medicine, Lund University, Malmö, Sweden

Twitter Katharina Beyer @beyer_katharina, Simone Scuderi @prostatePioneer, Riccardo Campi @Ric_Campi, Kirill Shiranov @endourologist, Muhammad Imran Omar @drimranomar, Alex Asiimwe @alexhabs and Sara J Maclennan @ SajMacLennan

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Contributors KB, LM, ML, AH, FB, SS, MM, MIO, SM, MJR, BF, EV, ZD, AA, JZ, SJM, LC, JN; ABr, ABj and MVH conceptualised designed the review. Abstracts and full texts were reviewed and data extracted by KB, LM, ML, AH, FB, SS, MM, RH, AR, RC, IG, KS, TvdB and SA. Authors resolved disagreement by discussion where necessary. The risk of bias was assessed by KB, LM, ML, AH, FB, SS, MM, RH, AR, RC, IG, KS and SD. The manuscript was drafted by KB, LM, ML, ABr, MVH and reviewed by KB, LM, ML, AH, FB, SS, MM, RH, AR, RC, IG, KS and SD. The manuscript was drafted by KB, LM, ML, ABr, MVH and reviewed by KB, LM, ML, AH, FB, SS, MM, RH, AR, RC, IG, KS, SD, TvdB, MG, GG, MIO, SM, MJR, BF, EV, ZD, AA, JZ, SJM, LC, JN, ABr, ABj and MVH. The whole project was supervised and guided by JZ, SJM, LC, JN, ABr, ABj and MVH. PIONEER Consotium acts as a gaurantor.

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ORCID iDs

Katharina Beyer http://orcid.org/0000-0002-8450-8850 Francesco Barletta http://orcid.org/0000-0003-3585-3217 Megan Molnar http://orcid.org/0000-0003-4937-2887 Riccardo Campi http://orcid.org/0000-0001-5237-0888 Muhammad Imran Omar http://orcid.org/0000-0002-1597-3126 Steven MacLennan http://orcid.org/0000-0002-2691-8421

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PRISMA 2020 Checklist

| Section and Topic | ltem # | Checklist item | Location where item is reported |
|-------------------------------|-----------|--|---------------------------------------|
| TITLE | 1 | | |
| Title | 1 | Identify the report as a systematic review. | p1 |
| ABSTRACT | 1 | | |
| Abstract | 2 | See the PRISMA 2020 for Abstracts checklist. | P2 |
| INTRODUCTION | 1 | | |
| Rationale | 3 | Describe the rationale for the review in the context of existing knowledge. | P3 |
| Objectives | 4 | Provide an explicit statement of the objective(s) or question(s) the review addresses. | P4 |
| METHODS | 1 | | |
| Eligibility criteria | 5 | Specify the inclusion and exclusion criteria for the review and how studies were grouped for the syntheses. | P5 |
| Information sources | 6 | Specify all databases, registers, websites, organisations, reference lists and other sources searched or consulted to identify studies. Specify the date when each source was last searched or consulted. | P5 |
| Search strategy | 7 | Present the full search strategies for all databases, registers and websites, including any filters and limits used. | P5 |
| Selection process | 8 | Specify the methods used to decide whether a study met the inclusion criteria of the review, including how many reviewers screened each record and each report retrieved, whether they worked independently, and if applicable, details of automation tools used in the process. | P5 |
| Data collection process | 9 | Specify the methods used to collect data from reports, including how many reviewers collected data from each report, whether they worked independently, any processes for obtaining or confirming data from study investigators, and if applicable, details of automation tools used in the process. | P5 |
| Data items | 10a | List and define all outcomes for which data were sought. Specify whether all results that were compatible with each outcome domain in each study were sought (e.g. for all measures, time points, analyses), and if not, the methods used to decide which results to collect. | P5 |
| | 10b | List and define all other variables for which data were sought (e.g. participant and intervention characteristics, funding sources). Describe any assumptions made about any missing or unclear information. | P5 |
| Study risk of bias assessment | 11 | Specify the methods used to assess risk of bias in the included studies, including details of the tool(s) used, how many reviewers assessed each study and whether they worked independently, and if applicable, details of automation tools used in the process. | P5 |
| Effect measures | 12 | Specify for each outcome the effect measure(s) (e.g. risk ratio, mean difference) used in the synthesis or presentation of results. | n/a |
| Synthesis methods | 13a | Describe the processes used to decide which studies were eligible for each synthesis (e.g. tabulating the study intervention characteristics and comparing against the planned groups for each synthesis (item #5)). | P5 |
| | 13b | Describe any methods required to prepare the data for presentation or synthesis, such as handling of missing summary statistics, or data conversions. | P5 |
| | 13c | Describe any methods used to tabulate or visually display results of individual studies and syntheses. | P5 |
| | 13d | Describe any methods used to synthesize results and provide a rationale for the choice(s). If meta-analysis was performed, describe the model(s), method(s) to identify the presence and extent of statistical heterogeneity, and software package(s) used. | P5 |
| | 13e | Describe any methods used to explore possible causes of heterogeneity among study results (e.g. subgroup analysis, meta-regression). | P5 |
| | 13f | Describe any sensitivity analyses conducted to assess robustness of the synthesized results. | P5 |
| Reporting bias assessment | 14 | Describe any methods used to assess risk of bias due to missing results in a synthesis (arising from reporting biases). | P6-8 |
| Certainty assessment | 15 | Describe any methods used to assess certainty (or confidence) in the body of evidence for an outcome. | n/a |

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PRISMA 2020 Checklist

| Section and Topic | Item # | Checklist item | Location where item is reported | | | | |
|--|-----------|--|---------------------------------------|--|--|--|--|
| RESULTS | 10- | Describe the vessiles of the second calenties avecage from the source of seconds identified in the second to the source of studies included in | DC 11 | | | | |
| Study selection | 16a | Describe the results of the search and selection process, from the number of records identified in the search to the number of studies included in the review, ideally using a flow diagram. | P6-11 | | | | |
| | 16b | Cite studies that might appear to meet the inclusion criteria, but which were excluded, and explain why they were excluded. | P6-11 | | | | |
| Study characteristics | 17 | Cite each included study and present its characteristics. | P6-11 | | | | |
| Risk of bias in studies | 18 | Present assessments of risk of bias for each included study. | | | | | |
| Results of individual studies | 19 | or all outcomes, present, for each study: (a) summary statistics for each group (where appropriate) and (b) an effect estimate and its precision I.g. confidence/credible interval), ideally using structured tables or plots. | | | | | |
| Results of | 20a | For each synthesis, briefly summarise the characteristics and risk of bias among contributing studies. | P6-11 | | | | |
| syntheses | 20b | Present results of all statistical syntheses conducted. If meta-analysis was done, present for each the summary estimate and its precision (e.g. confidence/credible interval) and measures of statistical heterogeneity. If comparing groups, describe the direction of the effect. | / | | | | |
| | 20c | Present results of all investigations of possible causes of heterogeneity among study results. | / | | | | |
| | 20d | Present results of all sensitivity analyses conducted to assess the robustness of the synthesized results. | / | | | | |
| Reporting biases | 21 | Present assessments of risk of bias due to missing results (arising from reporting biases) for each synthesis assessed. | / | | | | |
| Certainty of evidence | 22 | Present assessments of certainty (or confidence) in the body of evidence for each outcome assessed. | / | | | | |
| DISCUSSION | | | | | | | |
| Discussion | 23a | Provide a general interpretation of the results in the context of other evidence. | P12 | | | | |
| | 23b | Discuss any limitations of the evidence included in the review. | P15 | | | | |
| | 23c | Discuss any limitations of the review processes used. | P15 | | | | |
| | 23d | Discuss implications of the results for practice, policy, and future research. | P13-14 | | | | |
| OTHER INFORMA | TION | | | | | | |
| Registration and protocol | 24a | Provide registration information for the review, including register name and registration number, or state that the review was not registered. | Published with BMJ open | | | | |
| | 24b | Indicate where the review protocol can be accessed, or state that a protocol was not prepared. | / | | | | |
| | 24c | Describe and explain any amendments to information provided at registration or in the protocol. | / | | | | |
| Support | 25 | Describe sources of financial or non-financial support for the review, and the role of the funders or sponsors in the review. | End of the manuscript | | | | |
| Competing interests | 26 | Declare any competing interests of review authors. | In submission | | | | |
| Availability of data, code and other materials | 27 | Report which of the following are publicly available and where they can be found: template data collection forms; data extracted from included studies; data used for all analyses; analytic code; any other materials used in the review. | Methods | | | | |





PRISMA 2020 Checklist

From: Page MJ, McKenzie JE, Bossuyt PM, Boutron I, Hoffmann TC, Mulrow CD, et al. The PRISMA 2020 statement: an updated guideline for reporting systematic reviews. BMJ 2021;372:n71. doi: 10.1136/bmj.n71

For more information, visit: http://www.prisma-statement.org/

Supplementary material Table 1: Search strategy

Database: OVID Medline Epub Ahead of Print, In-Process & Other Non-Indexed Citations, Ovid MEDLINE(R) Daily and Ovid MEDLINE(R) 1946 to Present, Embase <1974 to 2020 January 28>, EBM Reviews - Cochrane Database of Systematic Reviews <2005 to January 21, 2020> Search Strategy:

1 exp *Prostatic Neoplasms/ (262435)

2 exp *prostate cancer/ (245472)

3 (prostat* adj2 (cancer* or carcinoma* or malignan* or tumor* or tumour* or neoplas* or adenocarcinoma* or adenoma*)).tw. (332251)

4 or/1-3 (366427)

5 ((diagnostic or prognos* or predict*) adj10 (biomarker or biomarkers or factor or factors)).tw,kw. (717487)

6 ((diagnostic or prognos* or predict*) adj10 (Oncotype Dx Prostate or Prolaris or Decipher or Decipher PORTOS or ProMark)).tw,kw. (458)

7 5 or 6 (717869)

8 4 and 7 (17456)

9 limit 8 to english language [Limit not valid in CDSR; records were retained] (16484)

10 limit 9 to yr="2014 -Current" (8417)

11 conference abstract.pt. or Congresses as Topic/ or Conference Review.pt. or "Journal: Conference Abstract".pt. (3815712)

12 10 not 11 (5902)

13 (exp animals/ or exp animal/ or exp nonhuman/ or exp animal experiment/ or animal model/ or animal tissue/ or non human/ or (rat or rats or mice or mouse or swine or porcine or murine or sheep or lambs or pigs or piglets or rabbit or rabbits or cat or cats or dog or dogs or cattle or bovine or monkey or monkeys or trout or marmoset\$1).ti.) not (humans/ or human/ or human experiment/ or (human* or men or women or patients or subjects).tw.) (10251935)

14 12 not 13 (5882)

15 note/ or editorial/ or letter/ or Comment/ or news/ or (note or editorial or letter or Comment or news).pt. (4565255)

16 14 not 15 (5811)

17 (child/ or Pediatrics/ or Adolescent/ or Infant/ or adolescence/ or newborn/ or (baby or babies or child or children or pediatric* or paediatric* or peadiatric* or infant* or infancy or neonat* or 18

19

20

21

22

16 not 17 (5794)

10 use coch (6) 19 or 20 (5794)

18 use ppez,oemezd (5788)

remove duplicates from 21 (3140)

newborn* or new born* or adolescen* or preschool or pre-school or toddler*).tw.) not (adult/ or

aged/ or (aged or adult* or elder* or senior* or men or women).tw.) (4146377)

Supplementary material Table 2. Multidisciplinary expert meeting

| Profession | Attendance |
|--------------------------------------|------------|
| Urologist | Accepted |
| Epidemiologist | Accepted |
| Urologist | Accepted |
| Epidemiologist/Pharma representative | Accepted |
| Pathologist | Accepted |
| Urologist | Accepted |
| Urologist | Accepted |
| Epidemiologist | Accepted |
| Methodologist | Accepted |
| Epidemiologist/Pharma representative | Accepted |
| Urologist | Accepted |
| Urologist/Methodologist | Accepted |
| Urologist | Accepted |
| Urologist | Accepted |
| Oncologist | Accepted |
| Pharma representative | Accepted |
| Pharma representative | Accepted |
| Statistician/ Pharma representative | Accepted |

Table 3: PROBAST overall assessment

| Criteria | Reaching and overall judgement of RoB | |
|---|--|--|
| All domains are rated low risk. | Paper was classified as low RoB and low Applicability. | |
| One or more domain was judged to be high risk of bias. | Paper was classified as high RoB and high Applicability. | |
| One or more domain was judged to be unclear risk of bias. | Paper was classified as unclear RoB and h igh Applicability. | |

Supplementary material Table 4: QUIPS scoring

| Score of 6 domains | Overall RoB | |
|---|----------------------------------|--|
| All domains were classified as having low | Paper was classified as low RoB | |
| RoB, or up to one moderate RoB. | | |
| One or more domains were classified as | Paper was classified as high RoB | |
| having high RoB, or ≥ 3 moderate RoB. | | |
| All papers in between. | Paper was classified as having | |
| | moderate RoB | |