

## STUDY PROTOCOL

# The ReIMAGINE prostate cancer risk study protocol: A prospective cohort study in men with a suspicion of prostate cancer who are referred onto an MRI-based diagnostic pathway with donation of tissue, blood and urine for biomarker analyses.

Teresa Marsden<sup>1,2\*</sup>, Neil McCartan<sup>1</sup>, Louise Brown<sup>3</sup>, Manuel Rodriguez-Justo<sup>4,5</sup>, Tom Syer<sup>6</sup>, Giorgio Brembilla<sup>6</sup>, Mieke Van Hemelrijck<sup>7</sup>, Ton Coolen<sup>8</sup>, Gerhardt Attard<sup>9</sup>, Shonit Punwani<sup>6</sup>, Caroline M. Moore<sup>1,2</sup>, Hashim U. Ahmed<sup>10,11‡</sup>, Mark Emberton<sup>1,2‡</sup>, on behalf of the ReIMAGINE Study Group<sup>¶</sup>

**1** UCL Division of Surgical & Interventional Sciences, University College London, London, United Kingdom, **2** Department of Urology, University College London Hospitals NHS Foundation Trust, London, United Kingdom, **3** MRC Clinical Trials Unit, University College London, London, United Kingdom, **4** Research Department of Pathology, University College London, London, United Kingdom, **5** Department of Pathology, University College London Hospitals NHS Foundation Trust, London, United Kingdom, **6** Centre for Medical Imaging, University College London, London, United Kingdom, **7** School of Cancer and Pharmaceutical Sciences, Kings College London, London, United Kingdom, **8** London Institute for Mathematical Sciences, London, United Kingdom, **9** Cancer Institute, University College London, London, United Kingdom, **10** Imperial Prostate, Division of Surgery, Department of Surgery and Cancer, Faculty of Medicine, Imperial College London, London, United Kingdom, **11** Imperial Urology, Imperial College Healthcare NHS Trust, London, United Kingdom

‡ HUA and ME are joint senior authors on this work.

¶ Membership of the ReIMAGINE Study Group is listed in the [S8 File](#).

\* [teresa.marsden@ucl.ac.uk](mailto:teresa.marsden@ucl.ac.uk)



## OPEN ACCESS

**Citation:** Marsden T, McCartan N, Brown L, Rodriguez-Justo M, Syer T, Brembilla G, et al. (2022) The ReIMAGINE prostate cancer risk study protocol: A prospective cohort study in men with a suspicion of prostate cancer who are referred onto an MRI-based diagnostic pathway with donation of tissue, blood and urine for biomarker analyses. PLoS ONE 17(2): e0259672. <https://doi.org/10.1371/journal.pone.0259672>

**Editor:** Henry Woo, Australian National University College of Health and Medicine, AUSTRALIA

**Received:** July 5, 2021

**Accepted:** October 24, 2021

**Published:** February 24, 2022

**Copyright:** © 2022 Marsden et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Funding:** The Medical Research Council, U.K. (MRC), grant number MR/R014043/1 and Cancer Research U.K. (CRUK).

**Competing interests:** I have read the journal's policy and the authors of this manuscript have the following competing interests: Hashim U Ahmed receives grant funding from Sonacare Inc., Shonit Biocorp and Boston commercial entities.

## Abstract

### Introduction

The ReIMAGINE Consortium was conceived to develop risk-stratification models that might incorporate the full range of novel prostate cancer (PCa) diagnostics (both commercial and academic).

### Methods

ReIMAGINE Risk is an ethics approved (19/LO/1128) multicentre, prospective, observational cohort study which will recruit 1000 treatment-naïve men undergoing a multi-parametric MRI (mpMRI) due to an elevated PSA ( $\leq 20$ ng/ml) or abnormal prostate examination who subsequently had a suspicious mpMRI (score  $\geq 3$ , stage  $\leq T3bN0M0$ ). Primary outcomes include the detection of  $\geq$  Gleason 7 PCa at baseline and time to clinical progression, metastasis and death. Baseline blood, urine, and biopsy cores for fresh prostate tissue samples (2 targeted and 1 non-targeted) will be biobanked for future analysis. High-resolution scanning

Hashim U Ahmed is awarded academic funds from the MRC, CRUK, The Wellcome Trust, The British Medical Association (BMA), The Urology Foundaiton (TUF), Prostate Cancer U.K. (PCUK) and Imperial Health. Mark Emberton serves as a consultant/educator/trainer to Sonacare Inc., Exact Imaging, Angiodynamics Inc and Profound Medical. Acknowledgements - Mark Emberton receives research support from the United Kingdom's National Institute of Health Research (NIHR) UCLH/UCL Biomedical Research Centre. Caroline M Moore receives funding from the Prostate Cancer UK, Movember, the Medical Research Council, Cancer Research UK and the NIHR. She receives fees for HIFU proctoring from SonaCare. She has received speaker fees from Astellas, and Janssen. She carries out research into photodynamic therapy supported by Spectracure. Shonit Punwani receives research support from the United Kingdom's National Institute of Health Research (NIHR) UCLH/UCL Biomedical Research Centre Ton Coolen receives funding from Cancer Research UK and is Director of Saddle Point Science Ltd. This does not alter our adherence to PLOS ONE policies on sharing data and materials.

**Abbreviations:** eCRF, Electronic Case Report Form; EU, European Union; FFPE, formalin-fixed paraffin-embedded; H&E, haematoxylin and eosin; HTA, Human Tissue Authority; MRI, magnetic resonance imaging; MTA, material transfer agreement; mpMRI, multiparametric magnetic resonance imaging; NHS, National Health Service; PCa, prostate cancer; PSA, prostate specific antigen; REC, Research Ethics Committee; SOP, Standard Operating Procedure; TRUS, Transrectal Ultrasound; UCL, University College London.

of pathology whole-slide imaging and MRI-DICOM images will be collected. Consortium partners will be granted access to data and biobanks to develop and validate biomarkers using correlation to mpMRI, biopsy-based disease status and long-term clinical outcomes.

## Results

Recruitment began in September 2019 ( $n = 533$ ). A first site opened in September 2019 ( $n = 296$ ), a second in November 2019 ( $n = 210$ ) and a third in December 2020 ( $n = 27$ ). Acceptance to the study has been 65% and a mean of 36.5ml(SD $\pm$ 10.0), 12.9ml(SD $\pm$ 3.7) and 2.8ml(SD $\pm$ 0.7) urine, plasma and serum donated for research, respectively. There are currently 4 academic and 15 commercial partners spanning imaging (~9 radiomics, artificial intelligence/machine learning), fluidic (~3 blood-based and ~2urine-based) and tissue-based (~1) biomarkers.

## Conclusion

The consortium will develop, or adjust, risk models for PCa, and provide a platform for evaluating the role of novel diagnostics in the era of pre-biopsy MRI and targeted biopsy.

## 1 Introduction

The prostate cancer (PCa) diagnostic pathway has undergone significant change in recent years and a number of guidelines now recommend multi-parametric MRI as the first line investigation for those with suspected organ-confined disease [1–3]. Before this, diagnosis and risk stratification was reliant upon clinical examination, PSA and systematic transrectal ultrasound-guided (TRUS) biopsy, but we are now aware how imprecise this diagnostic strategy was. Recent studies that used an exacting reference standard showed that over half of men with clinically significant prostate cancer were incorrectly diagnosed when exposed to TRUS-biopsy [4].

The incorporation of imaging into the diagnostic pathway has nearly doubled the probability of accurate baseline risk stratification compared to what went before. This is because biopsy sampling is now directed to areas of radiologically abnormal tissue with a high probability of clinically significant PCa rather than reliance on a systematic sampling approach which is arguably a somewhat random process [4]. The result is fewer missed clinically significant cancers, less over-diagnosis of clinically insignificant PCa and less harm from fewer men biopsied.

### 1.1 Aims and objectives

Current prostate cancer risk models (largely in the form of risk calculators) continue to inform treatment allocation but are based upon data collected during an era in which systematic TRUS-biopsy was the standard of care. When the histological outputs of modern-day diagnostics are applied to such models there is a danger of risk inflation as ‘risk’ is amplified (towards the truth). The result is that men’s expected risk might be exaggerated and the risk of over-treatment increases as a result [5, 6].

The ReIMAGINE consortium was conceived to address the need for new or adjusted prostate cancer risk stratification models to reflect MRI-directed diagnostic strategy. The ReIMAGINE consortium is a collaboration between several academic and commercial

partners to use diagnostics that span imaging, fluidic and tissue-based data to define novel, measurable and deeply phenotyped mpMRI disease cohorts, known as endotypes. Within the ReIMAGINE Prostate Cancer Risk study, the principal work-strand of the ReIMAGINE Consortium, inputs collected during MR-led diagnostics will be used to inform novel baseline risk-stratification systems and prognostic models.

The longitudinal component of the study will assess cancer progression, time to metastasis and prostate cancer related death through national healthcare data linkage. Deep phenotyping of measurable mpMRI disease endotypes and long-term follow up will permit the first insights into characteristics that define significant disease and redefine risk stratification in the mpMRI diagnostic era.

## 2 Design and methods: The ReIMAGINE prostate cancer risk protocol

### 2.1 Study design

ReIMAGINE Prostate Cancer Risk is a multi-centre, prospective, observational, longitudinal cohort study of patients referred to secondary care with a suspicion of prostate cancer (elevated PSA and/or abnormal digital rectal exam) or those who are having further tests for re-stratification of pre-existing low risk disease. The study represents level 1b evidence for diagnostic and prognostic studies [7]. Participants will be recruited from a number of high-volume NHS prostate cancer diagnostic centres in London, U.K. where pre-biopsy mpMRI followed by MRI-targeted prostate biopsies are established as standard of care.

Consenting men will be asked to donate research samples (blood, urine and fresh prostate tissue) for biomarker analysis at the time of their NHS standard of care prostate biopsies. Consent will also be sought for research access to pre-biopsy prostate mpMRI data and national medical records until death (should further funding be secured).

Biomarker analysis of donated samples will complete the cross-sectional component of the study. Participants exit the study following collection of tissue samples and revert to the NHS standard of care prostate cancer pathway. Standard of care pathology results are provided during a routine NHS clinic in accordance with local hospital protocol.

This manuscript reflects the current protocol version 2.1, 12 November 2019.

### 2.2 Patient and public involvement (PPI)

The ReIMAGINE project includes a work strand which focuses solely on patient and public involvement (PPI). The PPI committee consists of patients, and the wider public, who have been affected by, or have experience of, prostate cancer. The committee meet at least once every three months to inform prioritisation of research outputs, steer media outputs, and assist with development of study materials e.g. participant information videos. All patient facing documents are reviewed by a PPI representative or Chair of the PPI Sub-Committee, and its members.

The PPI team has been responsible for the development of the ReIMAGINE Consortium website (<https://www.reimagine-pca.org>). The website hosts a range of patient and media information resources relevant to each work strand of the consortium. Resources include details of the consortium directors, consortium partners, and links to participant and public information videos. Additional COVID-19 specific resources have been provided on the website since the study resumed recruitment in April 2020, and outline the steps taken to reduce the risk of COVID-19 exposure during a study visit.

## 2.3 Patient population

ReIMAGINE Risk aims to recruit 1000 men who are undergoing standard of care mpMRI directed biopsy following detection of an MRI lesion. Men with a PSA of less than or equal to 20 ng/ml who have undergone an mpMRI prostate during an NHS diagnostic work-up will be invited to participate in the study if a lesion scoring 3, 4 or 5 (radiological stage  $\leq$  T3b; clinical or radiological stage N0 and M0) is identified on mpMRI and they have accepted the need for a targeted and systematic biopsy (Fig 1). Either Likert or PIRADSV2.1 scoring will be acceptable. There must be no history of prior prostate cancer treatment (chemical, biological, ablative, surgical, radiotherapy) and no anti-androgen exposure in the preceding 6 months (5-alpha reductase inhibitors permitted). Full inclusion and exclusion criteria are outlined in Table 1.

The SPIRIT schedule of enrolment, interventions, and assessments is provided to outline the aforementioned study design.

## 2.4 Exposure variable

The main exposure variable is the deep phenotyping of the MRI-visible lesion. This will be achieved using additional targeted sampling of MR-identifiable lesions using either visual-estimation or image-fusion software, dependent on standard approach at each recruiting centre. As such, underlying inter-tumoral heterogeneity will be captured [8, 9]. A control sample collected from an area of unremarkable tissue on MRI (Likert / PIRADSV2.1 score 1 or 2) will further inform the characterisation of MR lesions.

## 2.5 Phase of study

ReIMAGINE Prostate Cancer Risk is currently recruiting at three London sites. The first site opened to recruitment in September 2019, a second centre in November 2019 and a third in December 2020. Recruitment was briefly halted between March and April 2020 due to the COVID-19 pandemic.

So far, 533 participants have been recruited across three sites. Recruitment is expected to complete in March 2022.

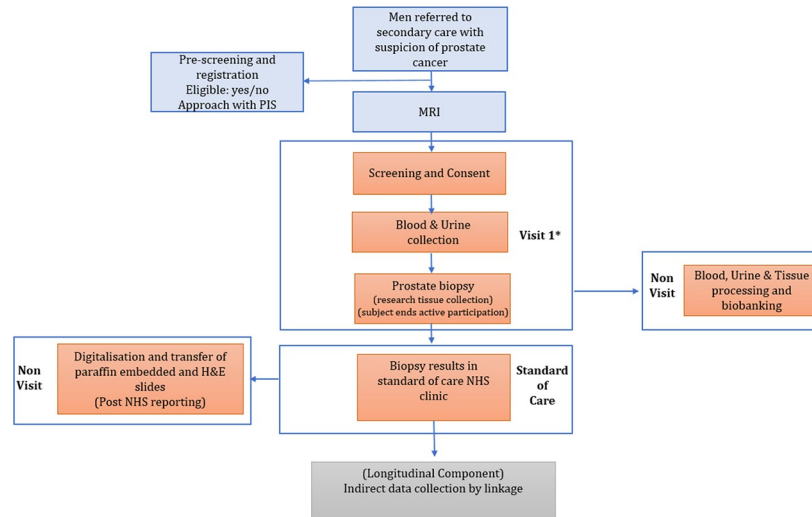
Acceptance to the study has been 65% and a mean of 36.5ml (SD $\pm$ 10.0), 12.9ml (SD $\pm$ 3.7) and 2.8ml (SD $\pm$ 0.7) urine, plasma and serum donated for research, respectively.

## 3 Study schedule

### 3.1 Screening and recruitment

Potential participants will be identified at the time of referral to a participating NHS centre, during a standard of care urology outpatient appointment or during screening of cancer waiting lists and theatre diaries. Participants will be approached in the first instance by either clinical staff or ReIMAGINE study staff (clinical trial practitioners). The first contact with eligible participants may be a telephone call or an invitation letter ahead of a standard of care prostate biopsy. Participating NHS centres may run a prostate cancer “one-stop” clinic whereby suspected cancer patients undergo biopsy on the day of baseline diagnostic mpMRI. Therefore, contact by telephone ahead of the visit is essential to maximise accrual. Potential participants will be offered a patient information leaflet to be sent electronically or by post ahead of their biopsy. All those approached or considered for recruitment will be recorded on the screening log.

During screening, and at the time of consent, patients will be assessed against the inclusion and exclusion criteria outlined in Table 1. Once eligibility is confirmed, written



	Study Duration			
	Screening	Visit 1	Non-visit	Indirect collection
TIMEPOINT**	-t <sub>1</sub>	0	3 – 6 weeks	t <sub>x</sub>
<b>ENROLMENT:</b>		X		
<i>Eligibility screen</i>	X			
<i>Informed consent</i>		X		
<i>Eligibility confirmation</i>	X	X		
<i>Medical History</i>	X	X		
<i>PSA</i>	X			
<b>INTERVENTIONS:</b>		X		
<i>Research blood</i>		X		
<i>Research urine</i>		X		
<i>Targeted biopsy</i>		X		
<i>FFPE tissue block and H&amp;E slide collection</i>			X	
<b>ASSESSMENTS:</b>				
<i>Standard of care biopsy results</i>			X	
<i>Adverse events review</i>		X		
<i>Concomitant medication review</i>	X			
<i>Healthcare data linkage</i>				X

**Fig 1. a.** ReIMAGINE Prostate Cancer Risk Study Design. ReIMAGINE Prostate Cancer Risk is a multi-centre, prospective, observational, longitudinal cohort study of patients referred to secondary care with a suspicion of prostate cancer (elevated PSA and/or abnormal digital rectal exam) or those who are having further tests for re-stratification of pre-existing low risk disease. Consenting men will be asked to donate research samples (blood, urine and fresh prostate tissue) for biomarker analysis at the time of their NHS standard of care prostate biopsies (Visit 1). Consent will also be sought for research access to pre-biopsy prostate mpMRI data and national medical records until death (should further funding be secured). Biomarker analysis of donated samples will complete the cross-sectional component of the study (both academic and commercial). Participants exit the study following collection of tissue samples and revert to the NHS standard of care prostate cancer pathway. Standard of care pathology results are provided during a routine NHS clinic in accordance with local hospital protocol. Access to standard of care diagnostic biopsy tissue will be sought once the participating centre has completed all necessary diagnostic evaluation. Formalin-fixed paraffin-embedded (FFPE) tissue blocks and haematoxylin and eosin (H&E) stained slides will be requested by ReIMAGINE staff and anonymised H&E slides will be scanned using high resolution scanners and uploaded to the ReIMAGINE digital pathology database. Consent will be sought to obtain long-term patient outcomes from national health records, such as the Office for National Statistics, NHS Digital, Public Health England and other applicable NHS information systems or national databases. Initially, data will be collected for three years after the last patient visit with the intention to collect data up to participant death once adequate funding is secured. This will complete the longitudinal component of the study. **b.** ReIMAGINE Prostate Cancer Risk Study SPIRIT schedule of enrolment, interventions, and assessments.

<https://doi.org/10.1371/journal.pone.0259672.g001>

informed consent will be sought. Eligible patients will be invited to consent to donation of blood, urine and three prostate tissue cores for biomarker analysis (to be collected at the time of standard of care biopsy, therefore no additional visits beyond the standard of care are required). Participants will also consent to donation of pseudonymised pre-biopsy mpMRI files.

**Table 1. ReIMAGINE risk study inclusion and exclusion criteria.**

<b>Inclusion Criteria</b>
Any man with a PSA 20ng/ml or less (value recorded <90 days before study entry)
Men who have undergone a prostate mpMRI as a standard NHS diagnostic work-up
mpMRI lesion (or area of suspicion) conforming to Likert / PIRADSV2.1 3, 4 or 5 undergoing a targeted and systematic transperineal biopsy
Radiological stage T3b or less
Clinical or radiological stage N0 and M0
No anti-androgen exposure in the preceding 6 months (5-alpha reductase inhibitors permitted)
No prior treatment for prostate cancer (chemical, biological, ablative, surgical, radiotherapy). Prior ADT permitted with adequate wash-out as described above (>6months)
Previous TURP is permitted
Willing and able to provide written informed consent
<b>Exclusion Criteria</b>
Men unable to donate tissue, blood or urine
Previous prostate cancer treatment
Previous prostate biopsy <12 months from date of the mpMRI scan used to assess study eligibility (scoring Likert / PIRADSV2.1 3, 4 or 5)

ReIMAGINE Prostate Cancer Risk is a multi-centre, prospective, observational, longitudinal cohort study of patients referred to secondary care with a suspicion of prostate cancer (elevated PSA and/or abnormal digital rectal exam) or those who are having further tests for re-stratification of pre-existing low risk disease. Men with a PSA of less than or equal to 20 ng/ml who have undergone an mpMRI prostate during an NHS diagnostic work-up will be invited to participate in the study if a lesion scoring 3, 4 or 5 (radiological stage  $\leq$  T3b; clinical or radiological stage N0 and M0) is identified on mpMRI and they have accepted the need for a targeted and systematic biopsy (Fig 1). Either Likert or PIRADSV2.1 scoring will be acceptable. There must be no history of prior prostate cancer treatment (chemical, biological, ablative, surgical, radiotherapy) and no anti-androgen exposure in the preceding 6 months (5-alpha reductase inhibitors permitted). Full inclusion and exclusion criteria are outlined in Table 1.

<https://doi.org/10.1371/journal.pone.0259672.t001>

Consent to access standard of care biopsy formalin-fixed paraffin-embedded (FFPE) tissue blocks and haematoxylin and eosin (H&E) slides will be requested once the holding NHS centre has fulfilled all diagnostic requirements. Agreement will also be sought for healthcare data linkage through NHS digital, Public Health England and other relevant bodies to inform longitudinal outcomes.

Recruiting sites will register fully eligible and consenting patients on a research database (Research Data Collection Service, REDCap) hosted securely by UCL. Project data will be periodically exported, after additional de-identification, from REDCap to a central integrated database (the ReIMAGINE Clinical Data Lake, CDL) hosted by Philips in the European Union (EU). Each participant will be allocated a unique study identification number. Identifiable data will be held on a separate REDCap database hosted only by UCL and used during the evaluation of longitudinal outcomes.

**3.1.1 Visit 1.** Following written informed consent, participant demographics, height, weight, medical history and medication history will be recorded. Biological samples (donated blood and first pass / early morning urine) will be collected on the day of standard of care biopsy. Three additional research core biopsies (two from the primary region of radiological interest and one control biopsy from an area of non-suspicious Likert / PIRADSv2.1 1 or 2 tissue within the contralateral lobe of the prostate) will be collected at the time of standard of care diagnostic biopsies (Fig 2). Samples will be processed and stored at a UCL laboratory until recruitment of all study participants is complete. Details of sample processing and storage are outlined in section 4.0. Once recruitment is complete, samples will be distributed to academic and commercial partners for analysis.

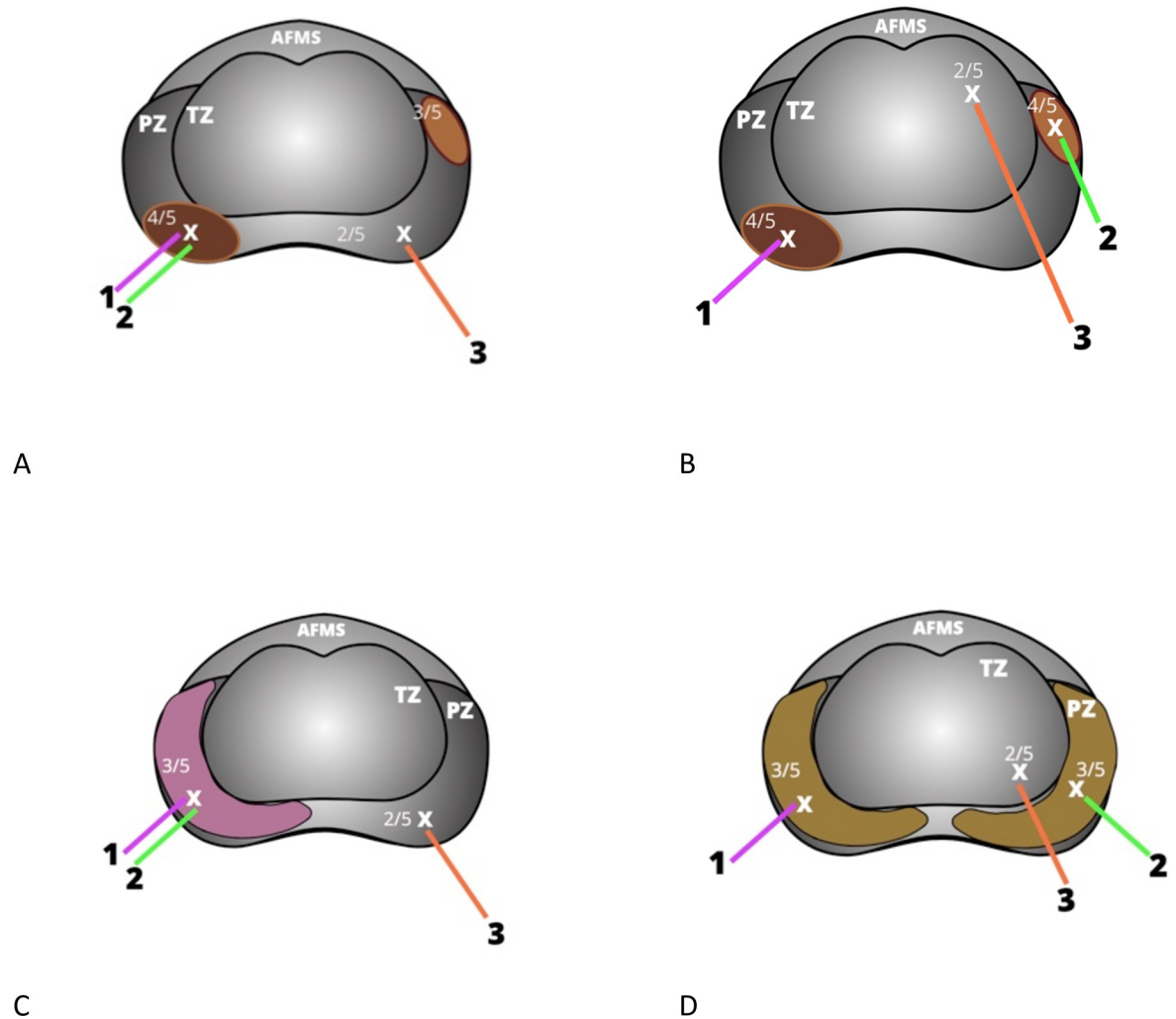
**3.1.2 Non-visit.** Access to standard of care diagnostic biopsy tissue will be sought once the participating centre has completed all necessary diagnostic evaluation. Formalin-fixed paraffin-embedded (FFPE) tissue blocks and haematoxylin and eosin (H&E) stained slides will be requested by ReIMAGINE staff and anonymised H&E slides will be scanned using high resolution scanners and uploaded to the ReIMAGINE digital pathology database. Slices from standard of care FFPE tissue blocks will be collected in line with study standard operating procedures before returning all slides and blocks to the relevant NHS pathology department.

**3.1.3 Analyses.** Blood components (white blood cells, plasma, serum) urine and tissue will undergo molecular analysis including genomic, mRNA expression and protein studies by academic and industry consortium partners. These will include, but are not limited to, next-generation sequencing, whole-genome micro-array expression profiling, immunohistochemistry and immunofluorescence and ELISA tests.

**3.1.4 Long-term healthcare data linkage.** Consent will be sought to obtain long-term patient outcomes from national health records, such as the Office for National Statistics, NHS Digital, Public Health England and other applicable NHS information systems or national databases. Initially, data will be collected for three years after the last patient visit with the intention to collect data up to participant death once adequate funding is secured. This will complete the longitudinal component of the study.

## 3.2 Definition of end of study

Following the collection of research samples (blood, urine and prostate tissue) men will revert to the NHS standard of care prostate cancer pathway at their centre. No further study visits are required. Men will undergo study follow up for 4 to 8 weeks after the biopsy for pathology results to be collected. Once complete, active data collection for the cross-sectional component of the study is complete. Longitudinal data collection through health data linkage will continue for three years and until death if adequate funding is secured.



**Fig 2. Location of research prostate tissue cores in ReIMAGINE prostate cancer risk.** A) Cores 1 and 2 are collected from the centre of the highest scoring lesion (denoted as Likert / PIRADSv2.1 score 4 in this figure) if there are no other lesions of the same score. The area of Likert / PIRADSv2.1 score 3 tissue in the left peripheral zone is not sampled for research. Core 3 is collected from an area of Likert / PIRADSv2.1 score 1 or 2 tissue (denoted as Likert / PIRADSv2.1 score 2 tissue in this figure). B) Where there is more than one lesion of the highest score, Core 1 is collected from the centre of the highest scoring lesion with greatest volume and Core 2 from the centre of the next largest lesion with the same score (both denoted as Likert / PIRADSv2.1 score 4 PZ lesions in this figure). Core 3 is collected from an area of Likert / PIRADSv2.1 score 1 or 2 tissue. C) If there is diffuse Likert / PIRADSv2.1 score 3 change to one peripheral zone (PZ) the cores should be collected from the centre of the area of score 3 change denoted on the radiology report (Cores 1 and 2). The third research core will be collected from an area of non-suspicious (Likert / PIRADSv2.1 score 1 or 2) tissue. D) If there is diffuse Likert / PIRADSv2.1 score 3 to the PZs bilaterally, the side of greater suspicion should be sampled first (Core 1), and a second core collected from the contralateral score 3 PZ (Core 2). The third research core will be collected from an area of non-suspicious (Likert / PIRADSv2.1 score 1 or 2) tissue.

<https://doi.org/10.1371/journal.pone.0259672.g002>

#### 4 Study interventions

ReIMAGINE prostate cancer risk will provide a platform for commercial and academic partners to run and calibrate novel biomarker tests within a cohort of suspected prostate cancer patients undergoing mpMRI targeted prostate biopsy. Once recruitment and sample collection is complete, specimens and pre-biopsy mpMRI data will be distributed to partner academic and commercial teams for analysis.



### 4.1 mpMRI databank

Multiparametric MR imaging will be performed as NHS standard of care [1, 10]. Reporting will be according to study SOPs and include Likert and PIRADSV2.1 scales documented within a structured electronic case report form (eCRF). Lesions, and the prostate itself, will be contoured and segmented prior to upload to the study image repository. Details of image processing and storage are outlined in Appendix I in [S1 File](#).

### 4.2 Blood sample collection

Between 50 and 100 ml of blood will be collected from each consenting study participant by a trained phlebotomist or ReIMAGINE Clinical Trial Practitioner during visit 1. Venepuncture and sample collection will be performed in accordance with the ReIMAGINE Risk Blood Sampling SOP. The required samples are outlined in [Table 2](#). A minimum blood donation of 50 ml is requested (up to a maximum of 100 ml). Whole blood samples will be processed by a Laboratory Technician or Clinical Trial Practitioner at a project-affiliated laboratory or an allocated laboratory area within the hospital site. Details of sample processing are outlined in [Appendix II in S2 File](#).

Any sample processed at clinical sites will be temporarily stored onsite at  $-80^{\circ}\text{C}$ . Following initial onsite processing, pseudonymised samples will be logged onto the study FreezerPro database. Samples will be transferred by courier to a study-affiliated UCL laboratory at regular intervals and stored at  $-80^{\circ}\text{C}$ .

### 4.3 Urine sample collection

A first pass or early morning urine sample will be collected in line with the study SOP by study practitioners during visit 1. Urine collection will always take place prior to prostate biopsy. The pseudonymised samples will be stored temporarily at local sites, transferred to a study-affiliated laboratory and logged within the study FreezerPro database. Both unspun and centrifuged urine will be stored within the study biobank. Spun samples will be centrifuged using a one-step process (3300g for 20 mins, with a break, at room temperature). Both centrifuged and unspun urine will be stored at  $-80^{\circ}\text{C}$  as will the residual cell pellet.

### 4.4 Prostate tissue collection

Consenting patients will donate three additional cores of prostate tissue at the time of their standard of care prostate biopsy ([Fig 2](#)). We do not anticipate that collection of three

**Table 2. Blood samples collected during the ReIMAGINE risk study.**

Order of draw (by collection tube)	Number of tubes to be collected	Volume per tube (ml)
EDTA	1	10
SST	1	8.5
Streck <sup>TM</sup>	1	10
EDTA	1	10
PAXgene <sup>®</sup> Blood RNA	1	2.5
EDTA	1	10
Extra EDTA (optional)	0–6	10

Between 50 and 100 ml of blood will be collected from each consenting study participant by a trained phlebotomist or ReIMAGINE Clinical Trial Practitioner during visit 1. Venepuncture and sample collection will be performed in accordance with the ReIMAGINE Risk Blood Sampling SOP. The required samples and order of draw is outlined in [Table 2](#).

<https://doi.org/10.1371/journal.pone.0259672.t002>

**Table 3. Side effect profile of a targeted transperineal prostate biopsy as stated in the ReIMAGINE prostate cancer risk study patient information sheet.**

Side effect	Proportion of men	Duration
Pain/discomfort in back passage	Almost all	Temporary for 1–2 days
Burning when passing urine	Almost all men	Self-resolving, 1–3 days
Bloody Urine	Almost all	Self-resolving, 1–7 days
Bloody Sperm	Almost all	Lasting up to 3 months
Poor erections	1–2 in 100	Temporary for 1–6 weeks
Infection of skin/urine	1–2 in 100	7 days with treatment with antibiotics
Infection of skin/urine needing admission and intravenous antibiotics	Less than 1 in 500	7–14 days with treatment with antibiotics. Rarely up to 28 days of antibiotics after leaving hospital.
Difficulty passing urine requiring catheter placement for up to a week. A catheter is a soft plastic tube placed into the bladder to drain urine.	1 in 100	3–7 days
Bruising of skin	Almost all	Self-resolving, 7–14 days
Bruising spread to scrotum	1 in 100	Self-resolving, 7–28 days

Eligible patients at recruiting sites will be provided with a study information leaflet at least 24 hours prior to consenting to study entry. The risk profile of targeted transperineal prostate biopsies is included in the study patient information leaflet and is shown in this figure.

<https://doi.org/10.1371/journal.pone.0259672.t003>

additional tissue cores will increase the likelihood of side effects or adverse events relative to standard of care biopsy alone. The risks and adverse outcomes of prostate biopsy quoted within the ReIMAGINE Risk patient information sheet are outlined in Table 3. Research cores will be targeted (visual-estimation or software fusion) to areas of interest identified on pre-biopsy mpMRI. Research cores will only be attempted after all required standard of care diagnostic biopsies have been collected.

All cores will be collected in line with the study Tissue Collection SOP. The first research core will be collected from the centre of the lesion with the highest Likert / PIRADSV2.1 score and greatest volume (lesion 1). The second sample will be collected from the centre of lesion 1 unless multiple lesions with the same Likert / PIRADSV2.1 score exist.

If multiple lesions with the same Likert / PIRADSV2.1 score exist, the second core should be collected from the next largest lesion with the same MRI score. Lesion volume will be determined by mpMRI reporting or MRI-US fusion software. If there is diffuse Likert / PIRADSV2.1 score 3 change to one peripheral zone (PZ) the core should be collected from the centre of the area denoted on the radiology report. If there is diffuse Likert / PIRADSV2.1 score 3 to the PZs bilaterally, the side of greater suspicion should be sampled first. Patients exhibiting diffuse change to the whole prostate should not be recruited.

The third research core will be collected from an area of non-suspicious (Likert / PIRADSV2.1 score 1 or 2) tissue. All biopsy cores should be collected from the centre of each targeted lesion. Table 4 summarises the location from which to collect each prostate biopsy core.

The location of each prostate core will be recorded within a study CRF (Fig 3) and each core placed within a colour coded pathology cassette before being placed within PAXgene® tissue fixative. Samples will be fixed in PAXgene® Tissue Fix reagent for at least 2–24 hours. Cores are then treated with PAXgene® Tissue Stabilizer concentrate and ethanol for at least 2 hours. All pseudonymised samples will be logged to the FreezerPro database and stored at -80°C following stabiliser treatment.

**Table 4. Prostate tissue sampling locations for the ReIMAGINE prostate cancer risk study.**

	Lesion to target	Area within lesion to target
<b>Core 1</b>	Lesion with the highest Likert / PIRADSV2.1 score and greatest volume (lesion 1).	Centre of lesion 1.
		If there is diffuse change throughout the PZs/TZs uni- or bi-laterally, target the centre of the area of greatest suspicion as defined by the reporting radiologist.
<b>Core 2</b>	Lesion 1, unless multiple lesions with the same Likert / PIRADSV2.1 score exist.	Centre of lesion 1 or centre of lesion 2 if multiple lesions with the same Likert / PIRADSV2.1 score exist.
	In which case, target the next largest lesion with the same Likert / PIRADSV2.1 score (lesion 2).	
<b>Core 3</b>	Collect from an area of non-suspicious (Likert / PIRADSV2.1 score 1 or 2) tissue.	Any non-suspicious area e.g. contralateral transition zone (TZ).

The first research core will be collected from the centre of the lesion with the highest Likert / PIRADSV2.1 score and greatest volume (lesion 1). The second sample will be collected from the centre of lesion 1 unless multiple lesions with the same Likert / PIRADSV2.1 score exist. If multiple lesions with the same Likert / PIRADSV2.1 score exist, the second core should be collected from the next largest lesion with the same MRI score. The third and final research core will be collected from an area of radiologically normal tissue (defined as Likert / PIRADSV2.1 1 or 2).

<https://doi.org/10.1371/journal.pone.0259672.t004>

All biological samples are processed and stored in line with the ReIMAGINE Risk Laboratory Manual. When indicated, samples will be transferred to consortium partners according to study standardised operating procedures and material transfer agreements (MTAs).

#### 4.5 Formalin-Fixed Paraffin-Embedded (FFPE) tissue blocks and Haematoxylin and Eosin (H&E) slide collection


In addition to the three prostate tissue cores collected purely for research purposes (Table 4), standard of care diagnostic pathology FFPE tissue blocks and H&E slides will be requested from each participating centre once all diagnostic evaluation is complete. Pseudonymised H&E slides will be scanned using the Hamamatsu Nanozoomer S360 high-throughput digital slide scanner at a resolution of 0.23  $\mu\text{m}/\text{pixel}$ . Digitalised, pseudonymised slides will be stored within a dedicated ReIMAGINE database held securely on UCL servers. Digitalised images, after additional de-identification, will be exported via the XNAT platform to the ReIMAGINE Clinical Data Lake (CDL) hosted by Philips in the EU to facilitate data sharing.

Digitalisation will facilitate computer-based assessment of morphological features and central pathology review. ReIMAGINE tissue technicians will also collect tissue slices from standard of care FFPE tissue blocks to provide additional tissue for analysis by consortium partners. All FFPE tissue blocks and H&E slides will then be returned to the relevant NHS pathology department. All processes will be conducted in accordance with study SOPs.


### 5 Outcomes

A lack of consensus regarding the definition of clinically significant prostate cancer remains. Much debate is focused upon the heterogenous cohort of intermediate risk, Gleason 7 disease. Recent studies have applied varied definitions of clinical significance including primary Gleason pattern 4, any grade of cancer core length  $\geq 6\text{mm}$  [4] or any core biopsy sample indicating Gleason 3 + 4 disease or greater [11].


By defining the primary outcome of the cross-sectional arm of ReIMAGINE Risk as the presence of any Gleason pattern 7 disease or greater we will gain insight into the baseline characteristics associated with intermediate risk disease which is likely to progress. Such



UCL



Imperial College  
London



NHS  
Royal Free London  
NHS Foundation Trust

Site (circle one):    **CXH**      UCLH      RFH

Samples collected by: \_\_\_\_\_      Date: \_\_\_\_/\_\_\_\_/\_\_\_\_

### ReIMAGINE TISSUE COLLECTION FORM

Study ID:         Patient initials:    Consented: Y  N

**Please circle 1 option only per line**

#### TISSUE


**TARGET 1**

Collected    Y     N

Time of collection: \_\_\_\_:\_\_\_\_

Lesion volume: \_\_\_\_ (ml)

Volume determined by:    MRI     MR-US



NOTES:


**TARGET 2**

Collected    Y     N

Time of collection: \_\_\_\_:\_\_\_\_

Lesion volume: \_\_\_\_ (ml)

Volume determined by:    MRI     MR-US



No. of cores collected:	Collected by:
_____	_____
No. of cores received:	Processed by:
_____	_____


**TARGET 3**

Collected    Y     N

Time of collection: \_\_\_\_:\_\_\_\_

Lesion volume: \_\_\_\_ (ml)

Volume determined by:    MRI     MR-US



**TIME IN TISSUE FIX REAGENT: \_\_\_\_:\_\_\_\_**

ReIMAGINE Prostate Cancer Risk  
Tissue Collection Form V3.3 dated 13Feb2020

**Fig 3. ReIMAGINE prostate cancer risk tissue collection form denoting the location of each donated prostate tissue research core.** Consenting patients will donate three research cores of prostate tissue at the time of their standard of care NHS prostate biopsy and the location of each will be recorded on a ReIMAGINE Prostate Cancer Risk tissue collection form. The first research core will be collected from the centre of the lesion with the highest Likert / PIRADSv2.1 score and greatest volume (lesion 1). The second sample will be collected from the centre of lesion 1 unless multiple lesions with the same Likert / PIRADSv2.1 score exist. If multiple lesions with the same Likert / PIRADSv2.1 score exist, the second core should be collected from the next largest lesion with the same MRI score. The third and final research core will be collected from an area of radiologically normal tissue (defined as Likert / PIRADSv2.1 1 or 2).

<https://doi.org/10.1371/journal.pone.0259672.g003>

information will permit greater stratification within this heterogeneous cohort and further elucidate the definition of clinical significance.

Longitudinal objectives achieved through long-term health data linkage include time to metastasis and / or prostate cancer related death. Longitudinal data will provide meaningful clinical correlation for novel biomarkers and risk models. Primary and secondary outcomes are presented in Table 5. Study outcomes will be published in peer-reviewed journal articles once available.

## 5.1 Mechanistic opportunity

The study provides further opportunity for discovery beyond the predefined primary and secondary outcomes. ReIMAGINE Risk biobanks will present an opportunity for computer-aided diagnostics, machine learning, calibration of commercial biomarkers (blood, urine, tissue and radiomic), the identification of MRI endotype molecular signatures and the stratification of MRI endotypes using clinical, radiomic and molecular inputs.

## 6 Statistical considerations

### 6.1 Sample size calculation

During the PRECISION study, 65% of patients with a PIRADS lesion  $\geq 3$  were found to have any cancer (Gleason score 3 + 3 or greater) detected on targeted prostate biopsy and 50% of targeted histology from PIRADS  $\geq 3$  lesions was considered clinically significant (Gleason score 3 + 4 or greater) [11]. Within the population of 1000 recruited participants during ReIMAGINE Risk, a conservative estimate will expect 60% of the cohort to have at least Gleason 3 + 3 disease on targeted histology of which two thirds will be considered clinically significant, defined as any Gleason pattern 7 or greater. Therefore, for the cross-sectional component of the analysis we expect 396 cases of clinically significant cancer and 604 cases of no cancer or clinically insignificant cancer.

Professional consensus is moving away from disease specific death rates towards metastatic progression as the endpoint with the greatest face, criterion and construct validity [12]. Time to metastasis and / or prostate cancer related death defines the primary outcome of the longitudinal component of the study. A cohort of 1000 men with an abnormal MRI should generate an event rate (time to metastases) of 10–20% at 10 years [13]. A conservative view would expect 100 events at 10 years.

**Table 5. Primary and secondary outcomes of the ReIMAGINE prostate cancer risk study.**

<b>Primary outcome: Cross-sectional component</b>
Presence of clinically significant prostate cancer confirmed on biopsy, defined as any Gleason pattern 7 or greater.
<b>Primary outcome: Longitudinal component</b>
Time to metastasis and/or prostate cancer related death.
<b>Secondary Outcomes</b>
Time to new prostate cancer in men without cancer at baseline.
Time to prostate cancer specific death and all-cause death in all men.
Time to cancer progression in men identified with prostate cancer at baseline.
<i>Progression is defined as the need for salvage therapy to the prostate following the primary treatment strategy, or, need for systemic therapy for prostate cancer treatment following failure of the primary treatment strategy.</i>

Primary and secondary outcomes of the ReIMAGINE Risk study include both cross-sectional and longitudinal components. This data will inform novel prognostic models for prostate cancer and provide a platform for the discovery and calibration of the full range of prostate cancer diagnostic tests (both commercial and academic).

<https://doi.org/10.1371/journal.pone.0259672.t005>

## 6.2 Statistical analysis

Statistical analysis will be performed according to pre-specified statistical methodology before the inspection of any outcome data. Logistic regression modelling will be used to analyse the factors associated with presence of clinically significant cancer. Cox regression modelling (or other appropriate survival methods if there is evidence of non-proportional hazards) will be used to analyse the factors associated with time to new cancers, cancer progression, metastasis and death. A list of pre-specified factors will be agreed and these will be systematically analysed using the aforementioned models in order to identify factors which are independently associated with each outcome.

In addition to the simple frequentist analysis approach, a more sophisticated analysis will be conducted. We will first apply the SaddlePoint Signature multivariate Bayesian regression pipeline, which was designed specifically for regression with high-dimensional covariates, in order to generate (i) optimal predictive covariate selections (suppressing false positive associations), (ii) quantitative characteristics of true associations (including covariate interactions, if relevant), (iii) personalised multivariate risk signatures, and (iv) quantifiers of the degree of outcome predictability. This will be done both for the full covariate set and for the individual covariate streams (e.g. imaging, genomic, phenotypic), allowing us to compare the outcome prediction power of different information sources, and their potential synergy. The most recent version of the pipeline (version 2.10.1), uses advanced overfitting correction protocols (based on the replica method) and robust internal validation processes. The risk scores for the separate streams will provide intuitive visualisations of the heterogeneity of the patient cohort. In a second stage we will apply Bayesian multivariate latent class analysis, using the source-specific risk signatures computed in the first stage as meta-covariates, and assess whether and how the cohort should be further stratified into subgroups with distinct associations and/or base hazard rates.

## 7 Ethical considerations

The study abides by the principles of the Declaration of Helsinki and received regulatory approval from the Regional Ethics Committee (London–Stanmore 19/LO/1128). ReIMAGINE Prostate Cancer Risk is published on clinicaltrials.gov (ClinicalTrials.gov identifier: [NCT04060589](https://clinicaltrials.gov/ct2/show/study/NCT04060589)).

## 8 Discussion and limitations

The prostate cancer diagnostic pathway has, until very recently, relied upon PSA-informed systematic biopsy, which is a method prone to considerable sampling error. Such inaccuracy has skewed risk calculators, rendered studies unrepresentative and quite probably misinformed healthcare policy. This represents a significant burden of diagnostic imprecision and accounts for a high burden of over-diagnosis and over-treatment as well as missed diagnoses of clinically important cancers. The use of pre-biopsy mpMRI corrects much of this inaccuracy and MRI positivity is closely correlated to tumour grade and volume—two surrogates of disease significance [4].

The ReIMAGINE consortium will exploit radiomic, molecular and targeted histological data to define novel and measurable image-based disease cohorts. Data from deeply phenotyped MRI cohorts and national healthcare records will provide inputs for progression modelling to more accurately predict disease status and outcomes. Exome and genome analysis of localised prostate and non-small-cell lung cancers suggest an association between intratumoural heterogeneity and poor prognosis [14, 15]. We therefore expect that integrated clinical, molecular (whether fluidic or tissue-based) and imaging analysis will provide more accurate

risk stratification than histological assessment—the current gold standard diagnostic test. Molecular profiling of advanced prostate cancers undertaken by other groups will provide a comparator for deeply phenotyped, organ-confined cancer cohorts within ReIMAGINE and add context to our results [16].

ReIMAGINE will also work with industry partners to facilitate the discovery, calibration and validation of fluidic biomarkers in blood and urine. Genome and exome analysis of plasma-extracted circulating tumour DNA has shown promise during the prediction of disease outcomes in advanced prostate cancer as well as the prediction of local disease recurrence in other cancer types [17, 18]. Plasma-extracted cell-free DNA is also used to guide the treatment of advanced non-small cell lung cancers (NSCLC) [19], whilst commercially available urinary exosome and methylome tests show great potential for predicting clinically significant prostate cancer more accurately than PSA [20, 21].

### 8.1 Limitations

The ReIMAGINE Risk protocol has some potential limitations. First, novel biomarker tests will be calibrated and validated against a population in whom a standard of care prostate biopsy has been deemed necessary following assessment with mpMRI. Biomarkers will therefore not be calibrated to patients who have not had MR imaging or in whom prostate mpMR imaging is unremarkable, however this could form part of further work. A prostate tissue sample from an area of Likert / PIRADsv2.1 score less than or equal to 2 will be collected from each participant to mitigate against the absence of a normal MRI cohort and act as a control. We also anticipate around one third of participants will have altogether benign prostate sampling, and will therefore form a further comparator group, albeit not representative of radiologically normal prostates as there is a requirement for an mpMRI Likert / PIRADsv2.1 score of 3 or greater to enter the study.

Second, many patients referred for secondary care assessment and therefore captured within the study population are referred because of a raised PSA level. Further work may be required to assess the applicability of findings to a cohort of men with a normal PSA. A number of patients within ReIMAGINE Risk will be recruited following participation within the ReIMAGINE screening study. ReIMAGINE screening has been designed to assess the feasibility of bi-parametric prostate MRI as a screening tool for prostate cancer. Patients may screen positive on the basis of either, or both, MRI positivity and raised PSA density. This cohort may broaden the spectrum of patients recruited to ReIMAGINE Risk to include a greater proportion of participants with a normal PSA.

Third, within participating centres, prostate cancer risk assessment and management decisions are already heavily directed by mpMRI findings. Longitudinal study outcomes may, therefore, fail to accurately reflect outcomes conferred by traditional risk stratification models and under-represent the need for disease re-stratification with contemporaneous MRI and targeted histological inputs. However, ReIMAGINE Risk represents the first attempt to redefine risk calculators during a new era of prostate cancer precision diagnostics.

## 9 Conclusion

As we transition to mpMRI-led prostate cancer diagnosis it is imperative that we redefine existing risk stratification models. Current risk models continue to inform practice but are based upon historic tissue archives and clinical cohorts in which TRUS biopsy was the baseline diagnostic test; they therefore cannot readily be applied to patients diagnosed by an mpMRI-targeted approach. ReIMAGINE Risk will deeply phenotype mpMRI disease cohorts and collect long-term oncological outcomes from national healthcare records. Study data will provide

image-based inputs for progression modelling and capture the full range of novel prostate cancer diagnostics.

## Supporting information

**S1 Checklist. SPIRIT 2013 checklist: Recommended items to address in a clinical trial protocol and related documents\*.**

(DOC)

**S1 File. Appendix I: mpMRI data management.**

(PDF)

**S2 File. Appendix II: Blood sample processing.**

(PDF)

**S3 File. Appendix III: Material and sample storage.**

(PDF)

**S4 File. Appendix IV: Discontinuation/withdrawal of participants.**

(PDF)

**S5 File. Appendix V: ReIMAGINE consortium structure, governance and work strands.**

(PDF)

**S6 File. Appendix VI: ReIMAGINE consortium partners (Academic and Commercial) at the time of publication.**

(PDF)

**S7 File. Appendix VII: Data management model.**

(PDF)

**S8 File. Appendix VIII: Acknowledgments.** The ReIMAGINE risk study group.

(DOCX)

**S9 File.**

(DOCX)

## Author Contributions

**Conceptualization:** Manuel Rodriguez-Justo, Mieke Van Hemelrijck, Gerhardt Attard, Shonit Punwani, Hashim U. Ahmed, Mark Emberton.

**Data curation:** Teresa Marsden, Neil McCartan, Tom Syer, Giorgio Brembilla, Ton Coolen, Hashim U. Ahmed.

**Formal analysis:** Louise Brown, Manuel Rodriguez-Justo, Mieke Van Hemelrijck, Gerhardt Attard, Shonit Punwani, Hashim U. Ahmed, Mark Emberton.

**Funding acquisition:** Gerhardt Attard, Shonit Punwani, Hashim U. Ahmed, Mark Emberton.

**Investigation:** Teresa Marsden, Manuel Rodriguez-Justo, Tom Syer, Giorgio Brembilla, Mieke Van Hemelrijck, Ton Coolen, Gerhardt Attard, Caroline M. Moore, Hashim U. Ahmed, Mark Emberton.

**Methodology:** Louise Brown, Manuel Rodriguez-Justo, Mieke Van Hemelrijck, Gerhardt Attard, Shonit Punwani, Hashim U. Ahmed, Mark Emberton.

**Project administration:** Neil McCartan.



**Resources:** Neil McCartan, Hashim U. Ahmed, Mark Emberton.

**Supervision:** Neil McCartan, Louise Brown, Manuel Rodriguez-Justo, Mieke Van Hemelrijck, Ton Coolen, Gerhardt Attard, Shonit Punwani, Caroline M. Moore, Hashim U. Ahmed, Mark Emberton.

**Visualization:** Hashim U. Ahmed, Mark Emberton.

**Writing – original draft:** Teresa Marsden, Neil McCartan, Gerhardt Attard, Shonit Punwani, Hashim U. Ahmed, Mark Emberton.

**Writing – review & editing:** Teresa Marsden, Neil McCartan, Louise Brown, Manuel Rodriguez-Justo, Tom Syer, Giorgio Brembilla, Mieke Van Hemelrijck, Ton Coolen, Gerhardt Attard, Shonit Punwani, Caroline M. Moore, Hashim U. Ahmed, Mark Emberton.

## References

1. Excellence NifHaC. NICE guideline [NG131]: Prostate cancer: diagnosis and management May 2019 [Available from: <https://www.nice.org.uk/guidance/ng131>
2. N. Mottet (Chair) PCV-c, R.C.N. van den Bergh, E. Briers (Patient Representative), M. De Santis, S. Fanti, S. Gillessen, et al. EAU Prostate Cancer Guideline 2019.
3. AUA: Pat F. Fulgham MDBR, MD; Jonathan N. Rubenstein, MD; Samir S. Taneja, MD; Peter R. Carroll, MD; Peter A. Pinto, MD; Marc A. Bjurlin, DO, MSc; et al. Standard Operating Procedure for Multiparametric Magnetic Resonance Imaging in the Diagnosis, Staging and Management of Prostate Cancer. A Collaborative Initiative by the American Urological Association and the Society of Abdominal Radiology Prostate Disease Focus Panel. May 2019 (Revised).
4. Ahmed HU, El-Shater Bosaily A, Brown LC, Gabe R, Kaplan R, Parmar MK, et al. Diagnostic accuracy of multi-parametric MRI and TRUS biopsy in prostate cancer (PROMIS): a paired validating confirmatory study. *Lancet*. 2017; 389(10071):815–22. [https://doi.org/10.1016/S0140-6736\(16\)32401-1](https://doi.org/10.1016/S0140-6736(16)32401-1) PMID: 28110982
5. Lomas DJ, Ahmed HU. All change in the prostate cancer diagnostic pathway. *Nat Rev Clin Oncol*. 2020; 17(6):372–81. <https://doi.org/10.1038/s41571-020-0332-z> PMID: 32112055
6. Vickers A, Carlsson SV, Cooperberg M. Routine Use of Magnetic Resonance Imaging for Early Detection of Prostate Cancer Is Not Justified by the Clinical Trial Evidence. *Eur Urol*. 2020.
7. Phillips B BC, Sackett D, et al Updated by Howick J. March 2009 [Available from: <http://www.cebm.net/oxford-centre-evidence-based-medicine-levels-evidence-march-2009/>
8. Dickinson L, Hu Y, Ahmed HU, Allen C, Kirkham AP, Emberton M, et al. Image-directed, tissue-preserving focal therapy of prostate cancer: a feasibility study of a novel deformable magnetic resonance-ultrasound (MR-US) registration system. *BJU Int*. 2013; 112(5):594–601. <https://doi.org/10.1111/bju.12223> PMID: 23819525
9. Hu Y, Kasivisvanathan V, Simmons LAM, Clarkson MJ, Thompson SA, Shah TT, et al. Development and Phantom Validation of a 3-D-Ultrasound-Guided System for Targeting MRI-Visible Lesions During Transrectal Prostate Biopsy. *IEEE Trans Biomed Eng*. 2017; 64(4):946–58. <https://doi.org/10.1109/TBME.2016.2582734> PMID: 27337710
10. Dickinson L, Ahmed HU, Allen C, Barentsz JO, Carey B, Futterer JJ, et al. Magnetic resonance imaging for the detection, localisation, and characterisation of prostate cancer: recommendations from a European consensus meeting. *Eur Urol*. 2011; 59(4):477–94. <https://doi.org/10.1016/j.eururo.2010.12.009> PMID: 21195536
11. Kasivisvanathan V, Rannikko AS, Borghi M, Panebianco V, Mynderse LA, Vaarala MH, et al. MRI-Targeted or Standard Biopsy for Prostate-Cancer Diagnosis. *N Engl J Med*. 2018; 378(19):1767–77. <https://doi.org/10.1056/NEJMoa1801993> PMID: 29552975
12. D'Amico AV. Active Surveillance Versus Treatment of Prostate Cancer: Should Metastasis Be the Primary End Point? *J Clin Oncol*. 2017; 35(15):1638–40. <https://doi.org/10.1200/JCO.2016.70.9527> PMID: 28240975
13. Johnston TJ, Shaw GL, Lamb AD, Parashar D, Greenberg D, Xiong T, et al. Mortality Among Men with Advanced Prostate Cancer Excluded from the ProtecT Trial. *Eur Urol*. 2017; 71(3):381–8. <https://doi.org/10.1016/j.eururo.2016.09.040> PMID: 27720537

14. Fraser M, Sabelnykova VY, Yamaguchi TN, Heisler LE, Livingstone J, Huang V, et al. Genomic hallmarks of localized, non-indolent prostate cancer. *Nature*. 2017; 541(7637):359–64. <https://doi.org/10.1038/nature20788> PMID: 28068672
15. Jamal-Hanjani M, Wilson GA, McGranahan N, Birkbak NJ, Watkins TBK, Veeriah S, et al. Tracking the Evolution of Non-Small-Cell Lung Cancer. *N Engl J Med*. 2017; 376(22):2109–21. <https://doi.org/10.1056/NEJMoa1616288> PMID: 28445112
16. James ND, Sydes MR, Clarke NW, Mason MD, Dearnaley DP, Spears MR, et al. Addition of docetaxel, zoledronic acid, or both to first-line long-term hormone therapy in prostate cancer (STAMPEDE): survival results from an adaptive, multiarm, multistage, platform randomised controlled trial. *Lancet*. 2016; 387(10024):1163–77. [https://doi.org/10.1016/S0140-6736\(15\)01037-5](https://doi.org/10.1016/S0140-6736(15)01037-5) PMID: 26719232
17. Annala M, Vandekerkhove G, Khalaf D, Taavitsainen S, Beja K, Warner EW, et al. Circulating Tumor DNA Genomics Correlate with Resistance to Abiraterone and Enzalutamide in Prostate Cancer. *Cancer Discov*. 2018; 8(4):444–57. <https://doi.org/10.1158/2159-8290.CD-17-0937> PMID: 29367197
18. Kidess E, Heirich K, Wiggan M, Vysotskaia V, Visser BC, Marziali A, et al. Mutation profiling of tumor DNA from plasma and tumor tissue of colorectal cancer patients with a novel, high-sensitivity multiplexed mutation detection platform. *Oncotarget*. 2015; 6(4):2549–61. <https://doi.org/10.18632/oncotarget.3041> PMID: 25575824
19. Jenkins S, Yang JC, Ramalingam SS, Yu K, Patel S, Weston S, et al. Plasma ctDNA Analysis for Detection of the EGFR T790M Mutation in Patients with Advanced Non-Small Cell Lung Cancer. *J Thorac Oncol*. 2017; 12(7):1061–70. <https://doi.org/10.1016/j.jtho.2017.04.003> PMID: 28428148
20. Donovan MJ, Noerholm M, Bentink S, Belzer S, Skog J, O'Neill V, et al. A molecular signature of PCA3 and ERG exosomal RNA from non-DRE urine is predictive of initial prostate biopsy result. *Prostate Cancer Prostatic Dis*. 2015; 18(4):370–5. <https://doi.org/10.1038/pcan.2015.40> PMID: 26345389
21. Leyten GH, Hessels D, Smit FP, Jannink SA, de Jong H, Melchers WJ, et al. Identification of a Candidate Gene Panel for the Early Diagnosis of Prostate Cancer. *Clin Cancer Res*. 2015; 21(13):3061–70. <https://doi.org/10.1158/1078-0432.CCR-14-3334> PMID: 25788493