

A Fully Automated, End-to-End Prostate MRI Workflow Solution Incorporating Dot, Ultrashort Biparametric Imaging and Deep-Learning-based Detection, Classification, and Reporting

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Introduction

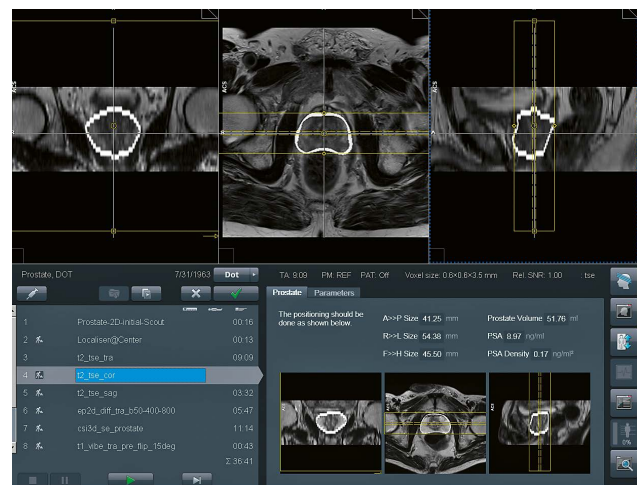
For more than a decade, magnetic resonance imaging (MRI) has been established as a powerful tool for prostate cancer diagnosis. The PROMIS study has demonstrated that prostate MRI is a suitable triage tool for biopsy-naïve men, reducing the number of unnecessary biopsies by a quarter while improving the detection of clinically significant cancer [1]. The PRECISION study randomized patients to either systematic biopsies or MRI; with no biopsy if MRI was negative, and targeted biopsy if MRI was positive. Targeted biopsies guided by MRI detected significantly more clinically significant cancers while reducing the number of clinically insignificant cancers [2]. Because of these findings, MRI for prostate cancer diagnosis has been integrated into established guidelines [3].

Increasing demand for prostate MRI examinations can be expected, as the incidence of prostate cancer increases with age and life-expectancy in developed countries is rising. Furthermore, prostate MRI has been discussed in the literature as a screening tool, similar to breast cancer screening [4]. However, several limitations need to be addressed in order to prepare for this increasing prostate MRI workload. Variation in MRI data acquisitions could be reduced [5]. Another limitation is the relatively long acquisition time of multiparametric MRI examinations (mpMRI) employing T2-weighted (T2w), diffusion-weighted imaging (DWI) and dynamic-contrast enhanced (DCE) MRI. Several studies have shown that an approach without DCE MRI, called biparametric MRI (bpMRI), yields comparable results to mpMRI of the prostate [6]. Potentially even more important topic is the varying interpretation performance

based on the expertise level. However, even among expert radiologists, agreement on prostate cancer classification based on established guidelines is imperfect [7, 8].

This all points to a clear need for

1. Efficient, reproducible, and robust data acquisition workflow
2. Optimized and fast sequence design
3. Automated detection, classification, and reporting workflows in prostate MRI examinations



1. Image acquisition using the Prostate Dot Engine¹ including automated prostate contour detection, prostate centering, field of view adaption and three-dimensional correction of spatial axes.

This is a chain of independent, yet highly interlinked stages. Well-registered and reformatted images with reproducible high image quality are a key prerequisite for optimal and reproducible artificial intelligence-based analyses.

In this article, we outline an end-to-end solution that addresses all the limitations above, incorporating day optimizing throughput (Dot), ultrashort bpMRI and deep-learning-based lesion detection, classification and reporting. We present two example cases using the proposed workflow in order to illustrate its feasibility.

Material and methods

Prostate Dot Engine

The Prostate Dot Engine¹ is a prototype software tool designed to provide a fast, robust, and standardized image acquisition workflow. After acquiring the Turbo-Spin Echo (TSE) scout, the Prostate Dot automatically centers the prostate in the field of view, adapts the size of the field of view and performs a three-dimensional correction of spatial axes. Slices can be aligned either strictly orthogonal or automatically defined by the orientation of the urethra, i.e., perpendicular to the urethra for the axial planes. Furthermore, the prostate is segmented for standardized volume assessment. After coil placement, the Dot workflow does not require further adaptations by technicians, and it allows interruptions and corrections of the scan process at any time. A screenshot of the Prostate Dot Engine can be found in Figure 1.

¹Work in progress: the application is currently under development and is not for sale in the U.S. and in other countries. Its future availability cannot be ensured.

Sequence specifications

The biparametric protocol consists of a T2-weighted turbo spin-echo (TSE) pulse sequence in axial, sagittal and coronal orientations and an improved single shot DWI EPI sequence (ZOOMit^{PRO}, Siemens Healthcare, Erlangen, Germany) with consecutive computation of the apparent diffusion coefficient. Unlike other DWI techniques, ZOOMit^{PRO} magnifies the prostate (in the phase-encoding direction) and is free of infolding artifacts. Either a smaller quadratic FOV or only a reduced FOV in the phase-encoding direction ('stripe') is excited (see Figure 2A). As there is no signal from the non-excited regions, only the small stripe needs to be encoded (see Figures 2B, C). That means the encoding time can be decreased while maintaining spatial resolution, or the spatial resolution can be increased, or a combination of the two. Furthermore, decreased encoding time reduces spatial distortion.

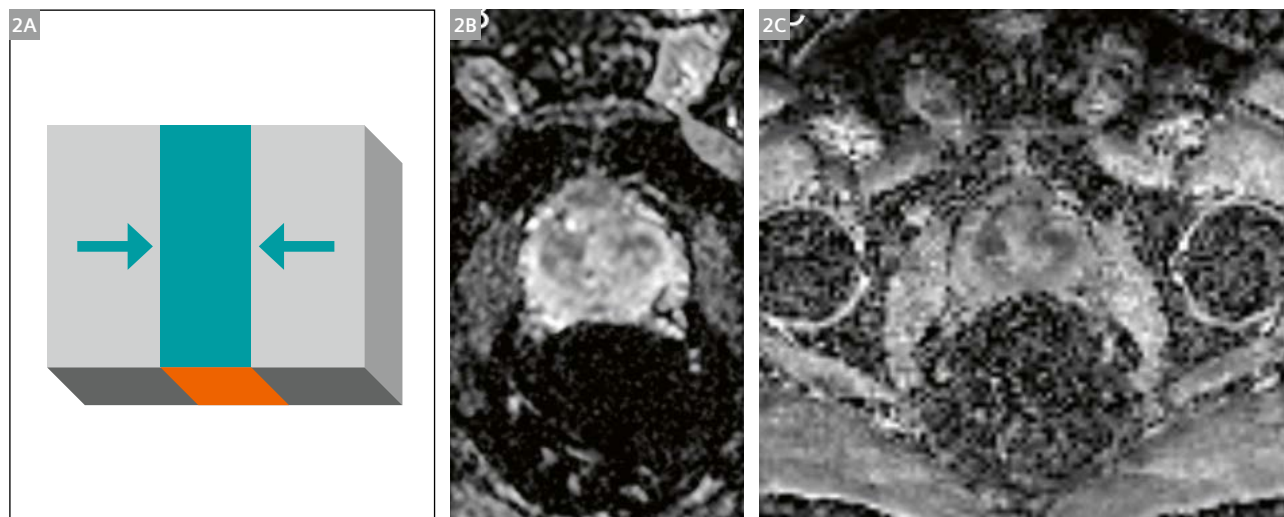
Prostate AI

The output of the Prostate Dot Engine goes into the AI prototype (Prostate AI¹, Siemens Healthcare, Erlangen, Germany) for fully automatic prostate lesion detection, classification and reporting.

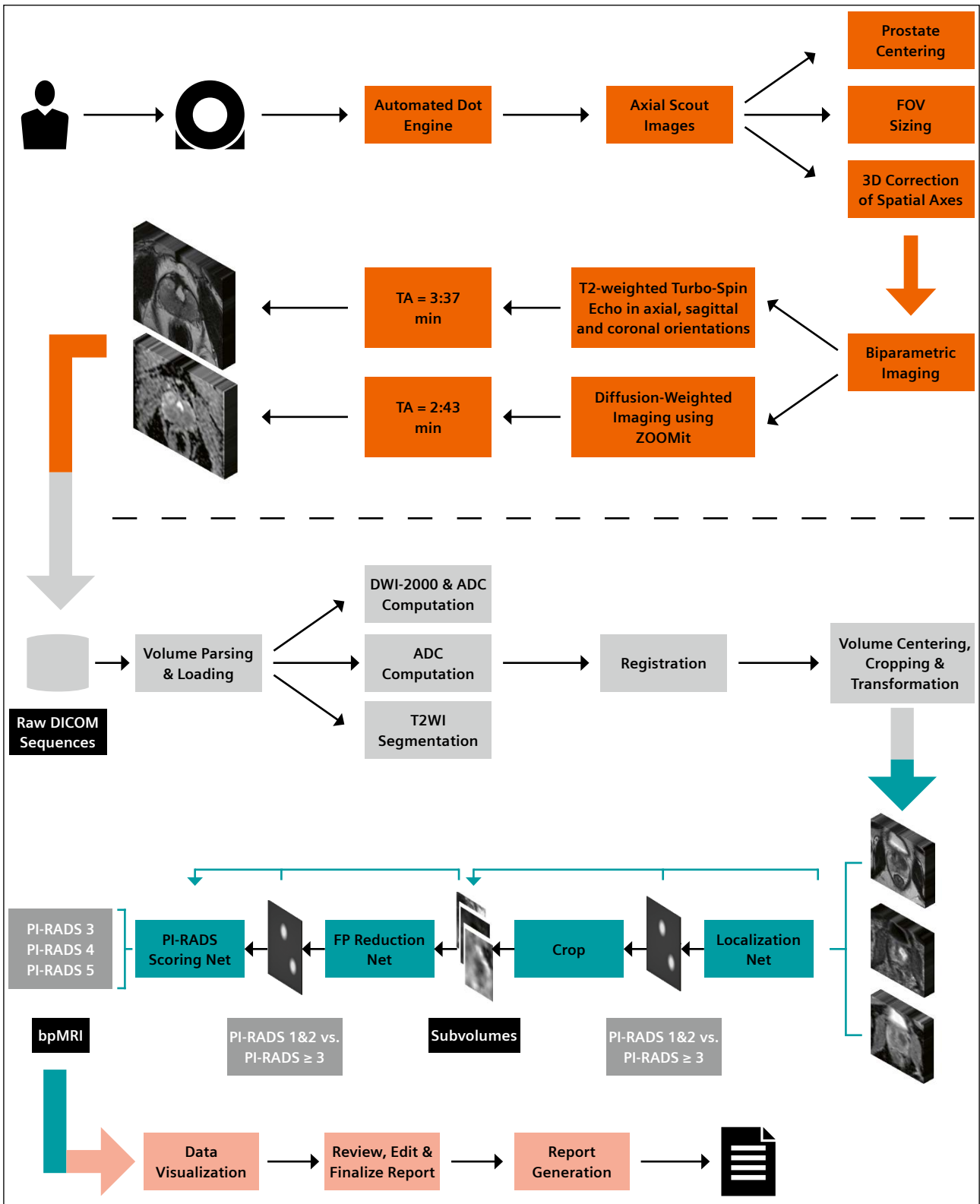
As illustrated in Figure 3, Prostate AI contains two parts:

1. A preprocessing pipeline
2. A component for lesion detection and classification, based on deep learning

The preprocessing pipeline takes the acquired bpMRI sequences and generates the required well-formatted and transformed data volumes. From the DWI series, a logarithmic extrapolation method is adopted to compute a new DWI volume with b -value of 2000 s/mm². This step

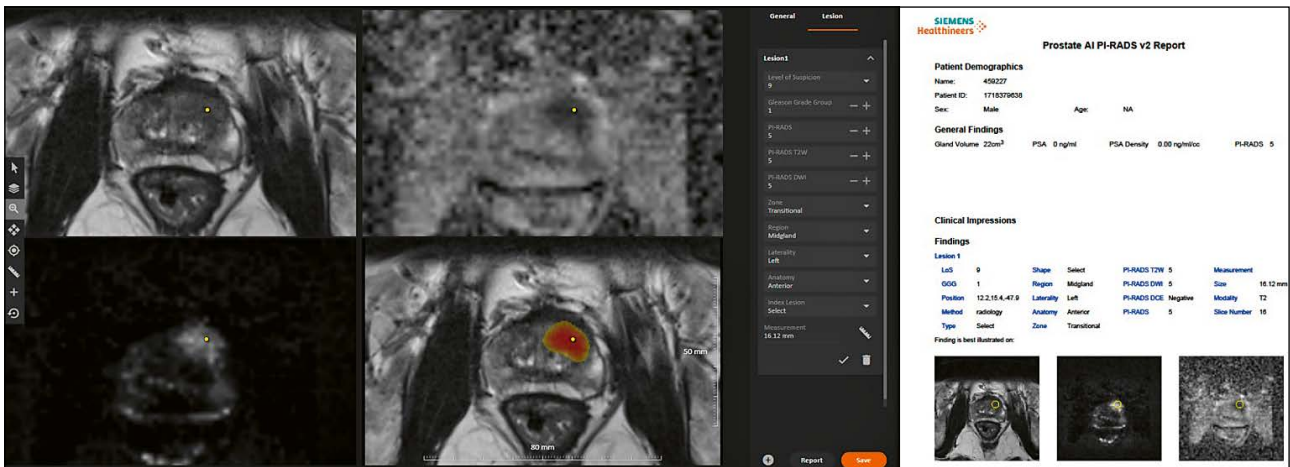


2 Single-shot DWI EPI sequence (ZOOMit^{PRO}) with image examples from one study object: **(2A)** reduced FOV in phase-encoding direction (blue stripe); **(2B)** resulting image in comparison to **(2C)** the conventional RESOLVE technique.



3 Image acquisition workflow using the automated Prostate Dot Engine and biparametric imaging (orange); deep learning architecture with preprocessing pipeline (gray); deep learning-based lesion detection and classification component (blue).

Dot = day optimizing throughput, FOV = field of view, 3D = three-dimensional, TA = time of acquisition, DICOM = Digital Imaging and Communications in Medicine, ADC = apparent diffusion coefficient, FP = false positive, PI-RADS = Prostate Imaging Reporting- and Data System



4 Data visualization platform with the T2w images, ADC map, and high *b*-value image as well as the T2w image overlaid with the AI-generated heatmap (in red and yellow). Prostate AI automatically detected the suspect lesion in the transition zone (TZ, yellow dot) and pre-populated all relevant information according to current PI-RADS guidelines. Next, a machine-readable report based on this information is generated.

can eliminate the *b*-value variances among the datasets and also improve lesion detection performance [10]. Also, apparent diffusion coefficient (ADC) maps are computed. Next, whole-organ gland segmentation is performed on the T2w volume using a learning-based method as presented in Yang et al. [11]. After segmentation, a rigid registration is conducted to align T2w and DWI images. The preprocessing pipeline can eliminate both geometric and intensity variances across sequences and patient studies.

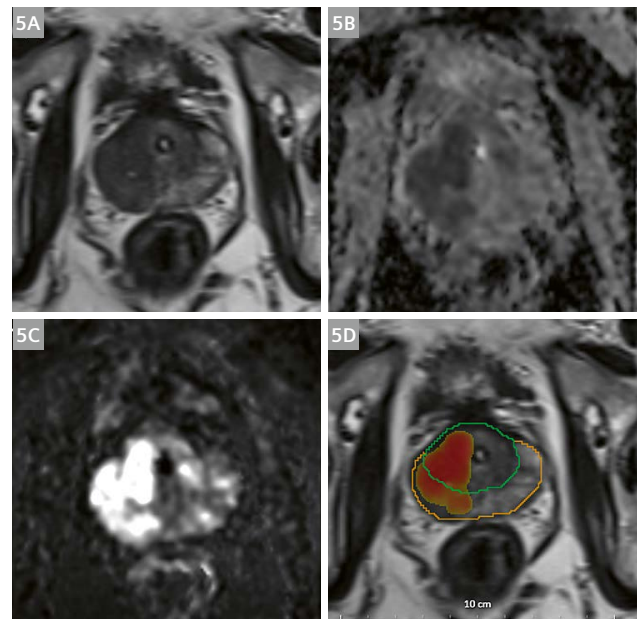
Prostate AI then automatically detects clinically relevant lesions and classifies each detected lesion according to PI-RADS categories. This is achieved by a sequence of coupled deep neural networks that are trained separately. First, a fully convolutional localization net is able to generate a semantic lesion candidate heatmap (see Figures 5 and 6); then a sub-volume-based false positive reduction net further improves detection accuracy by removing the false positives; finally another sub-volume-based PI-RADS scoring net stages the level of malignancy for each detection according to PI-RADS categories.

In a last step, Prostate AI displays the detection and classification results on a dedicated platform. As the ability of the interpreting radiologist to accept or reject AI-based findings has been identified as a prerequisite for adoption of these techniques [12], these capabilities have been implemented. The user is then able to create a machine-readable report with all relevant information for the referring physician (see Figure 4). This report can be sent to the local RIS/PACS system.

Cases

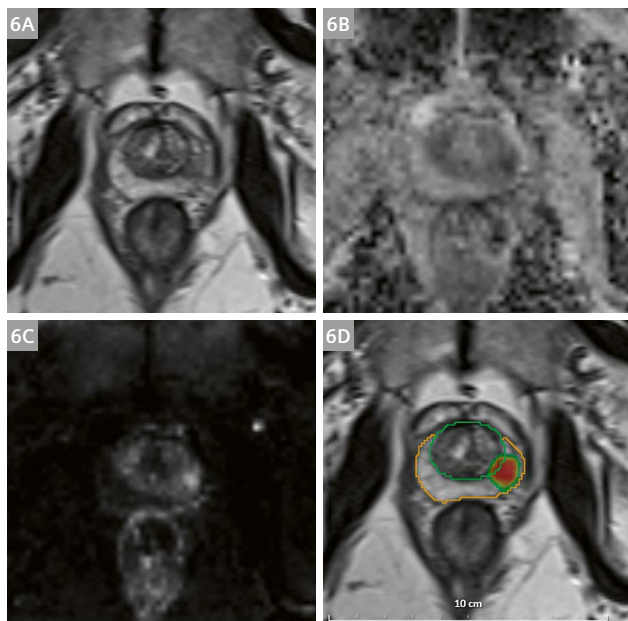
Case 1

Figures 5A-D demonstrate a lesion in the right midgland PZpl/PZa of a 62-year-old man, with a maximum diameter of 30.2 mm and a mean ADC-value of 758 $\mu\text{m}^2/\text{s}$. Prostate AI detected the lesion and assigned a PI-RADS 5 category. Biopsy results revealed a Gleason 4+3 = 7 pattern.



Case 2

Figures 6A-D demonstrate a lesion in the left apical PZpl of a 51-year-old man, with a maximum diameter of 10.2 mm and a mean ADC-value of $961 \mu\text{m}^2/\text{s}$. Prostate AI detected the lesion and assigned a PI-RADS 4 category. Biopsy results revealed a Gleason 3+3 = 6 pattern.



Conclusion

In this article, we outlined an end-to-end concept to allow a standardized workflow with a reproducible and fast data acquisition with optimized imaging sequences and an AI-empowered data analysis including automated detection, classification and reporting of suspicious lesions in biparametric prostate MRI examinations.

Reproducible and fast data acquisition concepts are not only contributing to a standardized reporting performed by human readers but would also help artificial intelligence-based solutions to reliably process input data. Preliminary results from a study conducted at the University of Innsbruck in Austria including 50 patients referred for a prostate MRI examination, compared the tilting angle of the auto-alignment of the Prostate Dot Engine against axes determined manually by an experienced radiologist, serving as the reference-standard. The investigators were able to show a mean \pm SD deviation of the tilting angle of 5.5 ± 4.4 degrees (Ch. Kremser, W. Judmaier, Med. Uni Innsbruck, unpublished results). However, to date,

there is no study investigating workflow differences, such as time-saving metrics, between Dot-guided and conventional, technician-guided workflows. Those studies are currently planned, and their results will contribute to reveal the value of Dot engines in clinical routine.

Concerning the use of abbreviated protocols consisting of T2-weighted and DWI only – so-called biparametric prostate MRI – several studies [6, 13, 14] have shown comparable results as obtained with conventional, mpMRI protocols including DCE-MRI. We added another component to our suggested workflow, that is performing DWI with the ZOOMit^{PRO}. As shown in Figure 2, ZOOMit^{PRO} uses a reduced FOV in the phase-encoding direction compared with either standard single shot DWI EPI or RESOLVE (REadout Segmentation Of Long Variable Echo trains). The resulting decreased acquisition time can be invested in a superior spatial resolution. Future studies are needed to systematically investigate differences between different types of DWI acquisition schemes compared to the ZOOMit^{PRO} technique.

The last component in our workflow is the use of AI-based lesion detection and classification. Schelb et al. [15] used the input from T2w sequences and DWI to train a deep learning algorithm (Unet) on the histopathological outcome, serving as ground truth. They were able to show that this algorithm achieved a similar performance to human readers using the PI-RADS assessment score. Cao et al. [16] used the input of mpMRI images to build a convolutional neural network trained on histopathological data and used this algorithm to detect suspicious lesions and to predict the Gleason score. The results were promising, with a high sensitivity for lesion detection – comparable to expert human readers – and a high classification performance with regards to clinically significant cancer. However, the usefulness of these algorithms needs to be proven in larger multi-reader, multi-case (MRMC) studies, systematically examining their influence on interpretation performance and speed, with and without those solutions.

We have identified a need to re-structure existing prostate MRI workflows, as patient or – in case of screening approaches – participant throughput is expected to increase. In our vision, current workflows need more reliable, reproducible and fast data acquisition steps. Furthermore, recent research has shown that deep learning algorithms can compete with human intelligence in prostate MRI reporting. We outlined a possible end-to-end solution and demonstrated its feasibility with two case examples. Future research will investigate what impact the individual components or the combination of those components will have on the future of prostate MRI.

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