

REVIEW

# A Systematic Review of Artificial Intelligence in Prostate Cancer

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<sup>1</sup>John P Hussman Institute for Human Genomics, Miller School of Medicine, University of Miami, Miami, FL, USA; <sup>2</sup>Department of Urology, Miller School of Medicine, University of Miami, Miami, FL, USA; <sup>3</sup>The Interdisciplinary Stem Cell Institute, Miller School of Medicine, University of Miami, Miami, FL, USA Abstract: The diagnosis and management of prostate cancer involves the interpretation of data from multiple modalities to aid in decision making. Tools like PSA levels, MRI guided biopsies, genomic biomarkers, and Gleason grading are used to diagnose, risk stratify, and then monitor patients during respective follow-ups. Nevertheless, diagnosis tracking and subsequent risk stratification often lend itself to significant subjectivity. Artificial intelligence (AI) can allow clinicians to recognize difficult relationships and manage enormous data sets, which is a task that is both extraordinarily difficult and time consuming for humans. By using AI algorithms and reducing the level of subjectivity, it is possible to use fewer resources while improving the overall efficiency and accuracy in prostate cancer diagnosis and management. Thus, this systematic review focuses on analyzing advancements in AI-based artificial neural networks (ANN) and their current role in prostate cancer diagnosis and management.

**Keywords:** prostate cancer, active surveillance, clinical trials, artificial intelligence

### Introduction

Prostate cancer is the most commonly diagnosed non-skin malignancy in men and second leading cause of mortality from cancer. Furthermore, it is estimated that 1 in 6 American men will be affected by this ailment during their lifetimes. Challenges faced during the course of prostate cancer management include increased volume of prostate biopsies and a shortage of urological pathologists, which puts a strain on diagnosis of prostate cancers. Additionally, the presence of a variability of pathological grading can result in an overtreatment or under treatment of prostate cancer.

Artificial Intelligence (AI) and machine learning in the healthcare is a new area of research which has recently gained a lot of interest. AI often makes use of artificial neural networks (ANN) that use statistical models that are directly inspired by and partially modeled on biological neural networks. They are capable of modeling and processing nonlinear relationships between inputs and outputs in parallel. Figure 1 shows a schematic of the structure of an ANN. In prostate cancer, the use of AI overall has shown to be beneficial to aid in a standardized pathological grading to assess prostate cancer stratification and treatment. Additionally, AI shows promise in automating the assessment of characterization and severity of prostate cancer based on image-based tasks including in histopathologic, MRI, and biomarker diagnosis. Furthermore, certain patients that are diagnosed with prostate cancer which is thought to be more indolent can continue with repeated forms of

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#### **Artificial Neural Network Architecture**

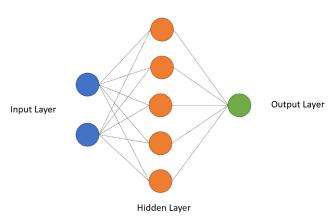


Figure I Schematic depicting the architecture of an artificial neural network.

surveillance including prostate biopsies, PSA, and other forms of digital testing through MRI or rectal examinations unless they experience any physiological side effects<sup>5</sup>. AI can help improve these forms of surveillance and will be amongst some of the essential tools to urological pathologists and to the field of urology as a whole as technology continues to improve and help patient prognosis over time. In this present study, we systematically reviewed the literature to determine how AI can be used to improve the diagnosis and management of prostate cancer.

### **Materials and Methods**

In this systematic review we used Pubmed, Embase, and Web of Science to identify and screen manuscripts pertaining to the use of machine learning and active surveillance in the diagnosis, staging, and treatment of prostate cancer. This review was conducted according to the Preferred Reporting Items for Systematic Review and Metaanalysis (PRISMA). The keywords used include: prostate biops\*) OR guided prostatic biops\*) OR ultrasound guided prostate biops\*) OR ultrasound guided prostatic biops\*) OR prostate biops\*) OR prostatic biops\*) OR gleason scor\*)) AND (((((artificial intelligence) OR deep learning) OR machine learning) OR artificial neural network) OR neural network)) AND (((clinical trial\*) OR active surveillance) OR active surveillance trial). Preliminary results showed 341 manuscripts. After duplicate removal, an exclusion criteria was applied to 283 manuscripts. This exclusion criteria is as follows: obviously not specific to the topic in question, case report, review, opinion, conference paper, commentary, survey, animal studies. After exclusion criteria was implemented on title and abstract, 81 full-text manuscripts were assessed for eligibility. After full-text assessment of eligibility, 19 manuscripts will be included in this systematic review. Figure 2 shows a flow diagram depicting the flow of information through the different phases of the literature search. A summary of included studies is presented in Table 1.

#### Literature Review

## Artificial Neural Network (ANN) and Prostate-Specific Antigens (PSA)

Identification of elevated PSA level is regarded as one of the most common clinical tool for diagnosis of prostate cancer. However, an elevated PSA level is present in several other benign prostatic conditions that can lead to unnecessary prognostic procedures. Generally, a PSA level greater than 4 ng/mL is considered for early detection of prostate cancer, but up to 20% of patients diagnosed with prostate cancer never reach that threshold. There has been a great need to diagnose prostate cancer in men with borderline PSA levels to improve health outcomes using new non-invasive methods.

Numerous studies have evaluated the utility of PSA and other clinical characteristics in the setting of an ANN in detecting prostate cancer and/or its progression. For example, Djavan et al developed an ANN to enhance the sensitivity and specificity of prostate cancer detection.<sup>8</sup> The study consisted of 1246 men with the primary aim of predicting the presence of prostate cancer or benign prostatic tissue in men with serum total PSA levels from 2.5 to 4 ng/mL and from 4 to 10 ng/mL compared to conventional tools. At 95% sensitivity, the ANN model for patients with total PSA levels from 4 to 10 ng/mL produced specificity, positive predictive value, negative predictive value, and receiver operator characteristic (ROC) curve area under the curve values that were superior to any of the other comparative parameters.8 A different study by Stojadinovic et al looked at various factors and determined that PSA density was the most decisive variable and showed the decision tree providing a net benefit compared to a logistic regression model. In this retrospective study, the model had a potential to reduce unnecessary biopsies without missing significant diagnoses.9

Finne et al developed an ANN using free PSA with the goal of eliminating unnecessary prognostic testing in false-positive PSA results.<sup>10</sup> The model was constructed using

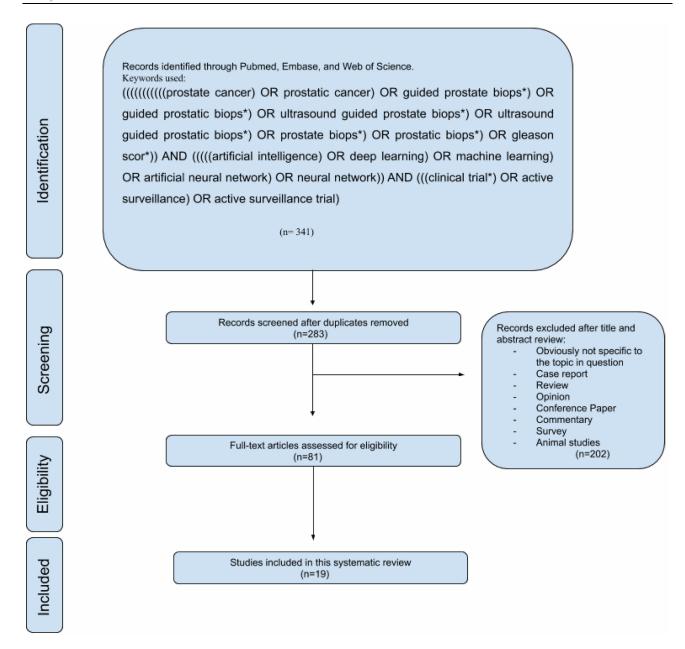


Figure 2 Flow diagram representing different phases of the systematic literature research according to PRISMA criteria.

Notes: PRISMA figure adapted from Liberati A, Altman D, Tetzlaff J, et al. The PRISMA statement for reporting systematic reviews and meta-analyses of studies that evaluate health care interventions: explanation and elaboration. Journal of clinical epidemiology. 2009;62(10). Creative Commons.<sup>43</sup>

data on total PSA, the proportion of free PSA, digital rectal examination (DRE), and prostate volume from 656 men with total serum PSA concentrations of 4 to 10 ng/mL in Finland. Results showed that, at a 95% sensitivity level, 19% of the false-positive PSA results could be eliminated by using the proportion of free PSA versus 24% with the logistic regression model and 33% with the ANN model. These findings suggest that an ANN method has value to be incorporated into prostate cancer screening to decrease the number of false-positive results. <sup>10</sup>

Ge et al developed a logistic regression and ANN model to aid in diagnosis of prostate cancer. They utilized predictors including age, percent free PSA, prostate volume, and PSA density from 586 men with prostate cancer confirmed by biopsy to train their algorithms. While they found no significant difference between the two models, they concluded that both models have high diagnostic validity and have potential to be included in practice to prevent unnecessary biopsies.<sup>11</sup>

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Table I Summary of All Articles Included in the Systematic Review

АІ Туре	AI Name	Summary	Author/ Reference
ANN + risk classification	PRODIGE	Proposed Umbrella Protocol that standardizes data and procedures to create a consistent dataset useful to elaborate Decision Support Systems. Thus this tool supports personalized decision making from multifactorial data sources.	Alitto et al <sup>38</sup>
ANN + MRI dx		Tested machine learning classifiers for transition zone and peripheral zone in MRI to classify prostate tumors with or without a Gleason 4 component. Classifiers trained within each zone had higher performance than the subjected option of pathologists.	Antonelli et al <sup>21</sup>
ANN + patient interaction	askMUSIC	Clinical registry to help patients interact with treatment decisions with similar characteristics.  Newly diagnosed patients can explore treatment options and compare their recommended treatments with other patients with similar treatments.	Auffenberg et al <sup>37</sup>
ANN + histopathologic dx		Automated image analysis using techniques similar to facial recognition to capture architectural differences between benign epithelium and various Gleason grades. This automated method showed concordance with trained pathologists and showed promise in differentiating G3+4 and G4+3 grades.	Bhele et al <sup>14</sup>
ANN + PSA dx		Statistical prediction of early prostate cancer by using PSA levels. The ANN accuracy level was higher than conventional PSA parameters and multivariate analysis.	Djavan et al <sup>8</sup>
ANN + MRI dx		Multiparametric MRI image processing combining the apparent diffusion coefficient and T2-weighted texture features. Results suggest texture features together with simple data augmentation offer reasonably accurate classification of Gleason patterns.	Fehr et al <sup>20</sup>
ANN + PSA dx		Multivariate algorithms based on clinical characteristics could reduce rate of false-positives in prostate screening more than free PSA alone. The logistic regression model showed higher accuracy and sensitivity when compared to free PSA showing promise to reduce the number of unnecessary prostate biopsies.	Finne et al <sup>9</sup>
ANN + PSA dx		Logistic regression and ANN diagnostic performance comparison. When aligned both algorithms showed no significant difference between the 2 results.	Ge et al <sup>10</sup>
ANN + biomarker dx		Predictive biomarkers for outcome in KI67 and DLX2. Both showed value to be able to inform clinical decision making in patients for active surveillance.	Green et al <sup>35</sup>
ANN + biomarker dx		Targeted proteomics to discover robust proteomic signatures for prostate cancer.  Computationally guided proteomics can be used to discover highly accurate non-invasive biomarkers.	Kim et al <sup>36</sup>
ANN + risk stratification		Using H&E slides, image processing used to be able to identify recurrence. Two level CNN has high accuracy when applied to 30 recurrent cases and 30 non-recurrent cases. This can possibly be applied to choose treatment options based on slides.	Kumar et al <sup>39</sup>
ANN + histopathologic dx		Using a polychotomous logistic regression model and ANN for predicting biopsy results.  Comparison of models showed no statistical difference between the two.	Lawrentschuk et al <sup>15</sup>
ANN + biomarker dx		Comparing statistical methods with ANN using conventional and experimental biomarkers.  Study showed the promise for both methods in evaluating prognostic markers and potential to use the technology to evaluate new markers.	Naguib et al <sup>40</sup>
ANN + PSA dx		Using ANN to predict recurrence based on free/total PSA, PSA density, and other clinical characteristics. The ANN found a pattern of prostate cancer with those patients that had a negative initial biopsy, and thus the ANN reduced unnecessary repeat biopsies.	Remzi et al <sup>17</sup>

(Continued)

Table I (Continued).

Al Type	AI Name	Summary	Author/ Reference
ANN + PSA dx		Used ANN to evaluate diagnostic value of % free PSA in men with total PSA levels between 2 and 20 $\mu$ g/L. The ANN showed enhanced accuracy when included with digital rectal examination and prostate volume measurements when compared to % free PSA alone.	Stephan et al <sup>11</sup>
ANN + PSA dx		Develop a classification and regression tree that could identify patients with significance prostate cancer based on patients with abnormal PSA, digital rectal examination findings, or both. The analysis showed provided net benefit when compared to a logistic regression model, PSA density, and biopsying all patients.	Stojadinovic et al <sup>9</sup>
ANN + histopathologic dx		The study evaluated the capacity of ANNs to assess prostate biopsies based on the presence, extent, and Gleason grade of the malignant tissue in comparison to experienced urological pathologists. Results showed that an ANN is able to be trained to detect and assess prostate biopsies with similar accuracy to urological pathologists.	Ström et al <sup>2</sup>
ANN + MRI dx		Classifier system for prediction of prostate cancer Gleason score using texture features of T2-weighted imaging in MRI images. Texture feature analysis showed good classification performance for GS in prostate cancer.	Toivonen et al <sup>23</sup>
ANN + histopathologic dx		Stratification of adenocarcinomas based on histological examination of tumor structure.  Complex patterns were stratified and then classified showing a reduction in intraobserver variability giving a better choice for patients to be recommended active surveillance.	Waliszewski et al <sup>14</sup>

Stephan et al developed an ANN utilizing percentage of free PSA (%fPSA) to identify men at increased risk of a positive prostate biopsy for cancer. The study enrolled 1188 men with benign prostates or prostate cancer with input data that included total PSA, %fPSA, patient age, prostate volume, and digital rectal examination (DRE) status. The use of the ANN in men with 2–10 µg/L total PSA enhanced the specificity of %fPSA by 20–22%. Thus, the use of an %fPSA-based ANN is able to enhance detection accuracy over %fPSA alone and prevent unnecessary biopsies. Together, there are multiple studies which reflect the relevance of using artificial intelligence-based ANN tools which could be effectively integrated with some of the vital tools commonly used in prostate cancer detection.

### Artificial Neural Network and Histopathologic Diagnosis of Prostate Cancer

The histopathological identification of prostatic adenocarcinoma is often essential for establishing a diagnosis of prostate cancer. However, sometimes histopathologic variability occurs due to inter- and intra-observer variability. This variability could impact the evaluation of biologic aggressiveness of prostate cancer and the identification of patients at high risk for progression. A solution to this

has been to introduce an artificial intelligence interface to accurately detect, localize and grade histopathologic slides. Early studies such as Bhele et al focused on detecting the difference in histology slides between Gleason score. Gleason grading allows the stratification of the aggressiveness of prostate cancer into low (grade group 1), intermediate (grade group 2 and 3) or high-risk group (grade group 4 and 5). Group 1 is designated by Gleason score (GS) 6, group 2 by GS7 or 3+4, group 3 by GS7 or 4+3, group 4 by GS8, and group 5 by GS9-10, respectively. The authors focused on detecting the difference between GS3 and GS4. 15 The model in this particular study was trained on 38 radical prostatectomy samples yielding 105 images. The tool reached a concordance on outlines of G3, G4, and benign epithelium of 67-81%. Thus, the AI model showed a good promise of being able to distinguish between 3+4 and 4+3 patients.

Using digitized slides from 1247 men, Ström et al trained ANNs to assess prostate biopsies.<sup>2</sup> The networks were evaluated by predicting the presence, extent, and Gleason grade of malignant tissue. The AI tool achieved an area under the ROC curve of 0.997 for distinguishing between benign and malignant biopsy cores on the independent test dataset and 0.986 on the external validation dataset. These results were comparable to the accuracy of

an international expert in prostate pathology.<sup>2</sup> An AI system such as this can provide a standardized grading method and pathology expertise in underserved areas.

by Lawrentschuk another study a polychotomous logistic regression (PR) model and an ANN was developed for predicting the biopsy results using 3025 men undergoing biopsy with PSA <10ng/ dL. 16 Clinical predictors of age, PSA, abnormal digital rectal examination, positive transrectal ultrasound (TRUS) and prostate volume were also considered during the development. It was suggested that an inclusion of additional predictors may improve the performance of the models; however, the ANN was unable to distinguish between the four biopsy outcomes that were utilized in their validation. The authors suggest that a more tempered enthusiasm should be used with ANN and that great care must be used in training them properly for effective application.<sup>17</sup>

In yet another study by Remzi et al, an ANN was developed to predict the presence of prostate cancer and predict the outcome in repeat biopsies. This model was trained using variables such as age, PSA, f/t PSA ratio, digital rectal examination findings, PSA velocity, and the transrectal ultrasound-guided variables of prostate volume, transition zone volume, PSAD, and PSA-TZ from 820 men with PSA levels between 4 and 10 ng/mL. This ANN showed a strong pattern of prediction for prostate cancer in patients who had an initial negative biopsy. Overall, the ANN achieved a 68% specificity and 95% sensitivity. Together, there are multiple studies which reflects the potential of AI ANNs to allow more accurate counseling of patients and overcome histopathologic variabilities.

### Artificial Neural Network and Magnetic Resonance Imaging (MRI) Diagnosis of Prostate Cancer

Magnetic resonance imaging (MRI) has been investigated as a modality for prostate cancer detection and determination of aggressiveness. <sup>19,20</sup> MR spectroscopic T2-weighted MR imaging, and apparent diffusion coefficient (ADC) have been valuable tools to assess prostate cancer, but their use still lacks consensus. <sup>21</sup> Using these tools, expert users can easily detect malignant tumors, but determining their aggressiveness using MRI with good reliability is more difficult. Thus, automatic classification using machine learning has been proposed as a solution to

provide more accurate and consistent results to aid clinicians' management. In the similar context, Fehr et al proposed a technique of utilizing machine learning-based automatic classification of prostate cancer aggressiveness by combining apparent diffusion coefficient (ADC) and T2-weighted MRI-based texture features. This technique was able to differentiate between the high GS (>7) and low GS6, as well as between 7 (3+4) and 7 (4+3) cancers. Furthermore, this model could distinguish between high and low Gleason grades with 93% accuracy for cancers occurring in both peripheral (PZ) and transition (TZ) zones and 92% for cancers occurring in the PZ alone. These results were far superior to using only ADC and suggest that this method can help provide reasonably accurate classification of Gleason patterns.

In addition, in another study by Antonelli et al, the authors used quantitative MRI and clinical features from 164 men to construct machine learning classifiers. <sup>22</sup> Following model validation, these classifiers were able to predict Gleason 4 in prostate tumors with greater accuracy than the three board-certified radiologists which participated in the study. It was suggested that these AI classifier tools could be useful to non-invasively detect the progression of tumors and aid in decisions regarding active surveillance programs. <sup>22</sup>

In yet another study by Toivonen et al, the authors developed machine learning tools to predict prostate cancer aggressiveness by using optimized high-quality MRI data sets.<sup>23</sup> A classifier system was created based on multiple texture features of high-quality T2 weighted images, DWI (diffusion weighted imaging), and T2 relaxation maps from 100 patients for prediction of PCa Gleason score dichotomized as 3+3 (low risk) vs >3+3 (high risk). Results suggested that texture feature analysis of DWI, post-processed using monoexponential and kurtosis models, and T2w demonstrated good classification performance for Gleason score of prostate cancer.<sup>23</sup> Together, these are multiple studies which reflects the potential of AI ANNs to allow more effective integration with modern surveillance tools such as MRI to effectively help in prostate cancer surveillance.

### Artificial Neural Network in Biomarker Diagnosis and Risk Stratification

Diagnosis and prognosis of prostate cancer has been guided by PSA level testing. However, concerns about absolute accuracy can sometimes lead to patients being

given invasive treatment options when active surveillance might provide better outcomes in these men.<sup>24</sup> Over the past 10 years, an avalanche of biomarkers have been identified and are put into clinical assays. 25-34 While several of these biomarkers have been studied and characterized, there is no standard overlap between all of these assays due to the function of each assay, and there is no absolutely perfect list of biomarkers to look at when predicting diagnosis and prognosis. Thus, it is important to be able to identify and evaluate any new biomarkers for their clinical significance in a way that is both meaningful and accurate. Therefore, ANNs can play an instrumental role in analyzing and validating the biomarkers. For example, one study suggested that Ki67 is an important marker of survival and disease progression.<sup>35</sup> To confirm this, Green et al created an ANN that was designed to validate Ki67 gene expression while comparing it to another potential candidate in DLX2.36 Univariate analysis showed that both Ki67 and DLX2 were significant in predictiveness of future metastases. Nevertheless, only 6.8% of prostate cancer patients have high expression of Ki67. Thus, this study showed that these 2 biomarkers could be used to identify candidates for targeted therapy only.<sup>36</sup>

In addition to gene expression, proteomics can also be useful when identifying potential biomarkers. For example, Kim et al developed a novel approach by combining targeted proteomics with computational biology to discover new potential proteomic signatures for prostate cancer. The study started with 133 differentially expressed proteins that were evaluated with synthetic peptides in a 74-patient cohort. Then they applied machine learning approaches to develop clinical predictive models using these candidates. Results showed that computationally guided proteomics can be used to discover novel non-invasive biomarkers. Overall, there are numerous studies which reflect the potential of AI ANNs to allow more effective identification and validation of biomarkers to aid in prostate cancer surveillance.

# Artificial Neural Network in Patient Interaction and Patient-Centered Treatment

Patients diagnosed with prostate cancer often remain confused as to the treatment options that are available to them. Thus, understanding how certain therapies are put into place may allow for ease of mind and improvements in patient satisfaction. Auffenberg et al developed a registry that used

ANN to better allow patients to take charge of their care. This registry, which is called askMUSIC, takes data from 45 urology practices within the Michigan Urological Surgery Improvement Collaborative (MUSIC). <sup>38</sup> This registry data is used to create a random forest machine learning model which could predict prostate cancer treatment options. Patients can go to askMUSIC website and interact with the registry data and predicted treatment to show therapy options to alleviate fear about a given therapy from the patient perspective.

In the similar context, the PRODIGE project by Alitto et al uses an Umbrella Protocol which focuses on standardization of data sharing. <sup>39</sup> Within this protocol, a standardized knowledge sharing process is implemented using semi-formal ontology representing clinical variables. This process can be adapted to use with machine learning or traditional statistics. The standardization of these techniques supports the multifactorial decision support systems (DSS) which can be seen as the basis for future patient-level support therapy decisions. Together, there are multiple studies which reflect the potential of AI ANNs to allow the development of effective patient-centric tools to help with educating the patients about the treatment options and disease progression/regression.

# Artificial Neural Network in Creating a Classification System for Prostate Cancer Risk Stratification

Current risk stratification as defined by the National Comprehensive Cancer Network (NCNN) guidelines relies on TNM (tumor, node, metastasis) score, Gleason grade, PSA level, and biopsy results.<sup>5</sup> These factors guide patients into risk groups which are stratified into 7 categories ranging from very low to very high. Based on these risk categories, physicians can guide patients into various personalized treatment options ranging from active surveillance, radical prostatectomy, radiation, and hormone therapy. While these risk categories are a good basis for stratification, none of the risk categories looks at the potential for recurrence. Thus, a better understanding of mechanisms behind recurrence could educate us to adjust the risk factors and therefore make the right choices for therapy.

Kumar et al focused on this recurrence prediction by creating 2 different convolutional neural networks (CNN) analyzing H&E images. 40 The first CNN detected individual nuclei while the second CNN classified the patches around the nuclear centers. There was a voting process which yielded probabilities of recurrence from the patients. The model was trained on 80 case/control pairs and then validated

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on 30 recurrent and 30 non-recurrent controls. The end yield was 0.81 AUC for accuracy. Thus, providing a deep learning tool that could look at recurrence in conjunction with other risk factors to come up with a new scoring system that could be more accurate than currently used.

In addition to recurrence, the current standards in identifying risk factors for prostate cancer often fails to take basic demographics into account. Nevertheless, a study by Naguib et al evaluated a combined approach of ANN and traditionally statistics known about various risk factors.<sup>41</sup> The neural network in this study was trained on conventional factors like age, stage, bone scan findings, grade, PSA and treatment along with 2 experimental markers of immunostaining for bcl-2 and p53. By considering conventional risk factors alone, the prediction models achieved only 60% accuracy in identifying the patients with condition. However, this efficiency was improved to 80% when the experimental markers were included. Thus, there was a significant improvement in accuracy of the model with an increase in the amount of data fed to the algorithm. This study also underscored the importance of including experimental markers for testing into building the networks as each piece of data could impact the model accuracy profoundly. Overall, there are multiple studies which reflect the potential of AI ANNs to allow the development of effective classification system for prostate cancer risk stratification.

### **Conclusion**

The use of artificial intelligence to manage medical problems has been discussed for quite some time. Nevertheless, it is only recently that technological advances have allowed us to make significant strides. AI can allow us to recognize difficult relationships and manage enormous data sets, which is a task that is both extraordinarily difficult and time consuming for humans. Moreover, the diagnosis and subsequent risk stratification that is required for active surveillance trials lends itself to enormous subjectivity. By using AI algorithms and reducing the level of subjectivity, it is possible to use fewer resources while improving the overall efficiency and accuracy in trials.

The economic burden of the management of localized prostate cancer must be acknowledged as well. The 10-year costs of managing low- and high-risk patients with prostate cancer have been estimated to be \$45,957 and \$188,928, respectively. However, a broader incorporation of AI can certainly help in mitigating this burden. Furthermore, AI makes it feasible to accomplish this while maintaining or perhaps even improving current

outcomes in active surveillance trials. Thus, it is essential to not only acknowledge but continue to improve the current pipelines to reduce costs, improve outcomes, and drive the future of urologic oncology forward as a whole.

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### **Disclosure**

The authors report no conflicts of interest for this work. Declaration of interests: the authors declare no competing interests.

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