

## BPH and prostate cancer risk

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### ABSTRACT

**Introduction:** With the exclusion of non-melanomatous skin malignancy, prostate cancer (PCa) is the second most prevalent cancer in men globally. It has been reported that the majority of men will develop benign prostatic hyperplasia (BPH) by the time they reach their 60s. Together, these prostatic diseases have a significant morbidity and mortality affecting over a billion men throughout the world. The risk of developing prostate cancer of men suffering BPH is one that has resulted in a healthy debate amongst the urological community. Here, we try to address this conundrum with clinical and basic science evidence.

**Materials and Methods:** Data from an online search and contemporary data presented at international urological congresses was reviewed.

**Results:** BPH and PCa can be linked together at a molecular and cellular level on genetic, hormonal, and inflammatory platforms suggesting that these prostatic diseases have common pathophysiological driving factors. Epidemiological studies are weighted towards the presence of BPH having a greater risk for a man to develop PCa in his lifetime; however, a conclusion of causality cannot be confidently stated.

**Conclusion:** The future workload healthcare practitioners will face regarding BPH, and PCa will substantially increase. Further basic science and large epidemiological studies using a global cohort of men are required prior to the urological community confidently counseling their patients with BPH with regards to their PCa risk.

**Key words:** Benign prostatic hyperplasia; epidemiology; epigenetics; genetics; inflammation; prostate cancer

### INTRODUCTION

The lifetime risk of a man developing histologically confirmed benign prostatic hyperplasia (BPH) has been reported to be 50% in those in the 51-60 years age group and increasing to 70% in the 61-70 years age group.<sup>[1]</sup> With the exclusion of non-melanomatous skin malignancy, prostate cancer (PCa) is the second most prevalent cancer in men globally. The International Agency for Research on Cancer reported that PCa accounted for 14% of cancers diagnosed in men and

over 900,000 cases were diagnosed throughout the world in 2008 alone.

The exact etiology of both these prostatic diseases has yet to be fully elucidated. Anatomically, BPH arises from the transition zone of the prostate gland. PCa is an adenocarcinoma, which classically arises from epithelial cells located in the peripheral zone of the prostate gland alongside a small percentage arising from those cells that are located in the transition zone. In only 20% of cases do BPH and PCa co-exist in the same prostatic zone.<sup>[2]</sup> In combination, both these diseases have a major influence on the morbidity and mortality in millions of men throughout the world. Advances in healthcare quality and provisions globally and the resultant increase in worldwide life-expectancy will ultimately lead to a significant percentage of men that will face either BPH or PCa individually or in combination. Furthermore, with the advent of more screening and disease detection strategies that have been adopted in national healthcare commitments throughout the world, the incidence and pick-up rates of these diseases will naturally increase in the younger population of men.

Despite both these diseases being well established histologically, a definitive link between them has proven

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to be one that leads to a healthy debate and controversy within the urological community. Both diseases have striking similarities with regards to androgen-dependence driving their pathophysiology, inflammatory components which contribute to the development of the diseases, and shared genetic and epigenetic alterations. However, a causal relationship between these diseases has not been established.

In this review, we address the question 'Is the presence of BPH a risk factor for the development of prostate cancer?' by using evidence ranging from basic science, genetic and major epidemiological studies conducted on this topic. The PubMed® was searched for relevant articles and, in addition, we examined secondary sources from the reference section of utilized articles as well as data presented at recent urological congresses.

### *Epidemiological association of BPH and PCa*

Historical autopsy data has shown that 83.3% of PCa arises with the concomitant presence of BPH.<sup>[3]</sup> Chokkalingam *et al.* reported the first large-scale population study over a 26-year period in men from the Swedish health database ( $n = 86,626$ ).<sup>[4]</sup> Men were clinically diagnosed with BPH either with no surgical intervention or surgical intervention in the form of a transvesical adenectomy or transurethral prostatectomy. It was concluded that in comparison to the general population, men with BPH experienced little if any increased risk of prostate carcinoma (2% after a 10-year follow-up). Patients who underwent a transvesicular adenectomy had a significantly (22%) lower incidence and a 23% lower mortality after the first 5 years of follow-up, and those who underwent a transurethral resection had a significantly higher incidence at 10% and a 17% lower mortality associated with PCa. Those with BPH who received no surgical intervention had significantly increased incidence and mortality associated with PCa at 18% and 77%, respectively. This study did demonstrate a mixed outcome of incidence and mortality of BPH patients in relation to PCa, and the authors correctly highlighted that the differences in statistics may be from underlying health factors creating bias for surgical treatment choice. Furthermore, all patients evaluated in this study were assessed in the pre-PSA era where two issues arise. First, the risk of prostate cancer-specific mortality associated with a finding of BPH will have been significantly higher. Second, the identification of PCa is made at an earlier stage during the current PSA-era in comparison to the pre-PSA era; therefore, the rate of discovering a new incidence of PCa from the surgical treatment of BPH has substantially decreased.<sup>[5]</sup>

The biggest epidemiological study to date to assess the risk of a man developing PCa following the diagnosis of BPH was reported by Orsted *et al.*,<sup>[6]</sup> in 2011. Using population-based databases, they evaluated this conundrum in the entire Danish male population over a 27-year period ( $n = 3,009,258$ ) between 1980 and 2007. The multivariate, adjusted hazard

ratios (HR) for PCa incidence/mortality was 2.22/2 (95% confidence interval, 2.13-2.31/1.91-2.08) amongst all men hospitalized for BPH compared to the general population control, respectively. This figure increased to 3.26/7.85 (95% CI; 3.03-3.5/7.40-8.32) when men who were surgically treated for BPH were taken into consideration versus the control group. Despite the authors' concluding that BPH was associated with a two-three-fold increased risk of the development of PCa along with a two to eight-fold increased risk of mortality, they stressed that this data should not be interpreted to infer causality. Limitations of this study included that the diagnosis of BPH was not a histological one but clinical, the relatively homogenous nature of this population group, and the issue that the study included use of patients in the pre-PSA era.

The association of BPH and PCa has also been demonstrated in men with an ethnic low-risk of PCa. Asian men who were diagnosed with prostate cancer over a 10-year period and all of whom were over the age of 50 years had a strong association with prostatitis and/or BPH.<sup>[7]</sup> There was a higher odds ratio (OR) for prostate cancer associated with BPH (26.2, 95% confidence interval (CI) 20.8-33.0) than with prostatitis (10.5, 95% CI = 3.36-32.7).

Despite large-scale studies demonstrating the epidemiological link between PCa and BPH, [Table 1], studies contradicting this finding have concluded that there is no association between these diseases. Schenk *et al.* assessed 1,549 men with BPH over a 20-year period between 1993 and 2003 and concluded that the presence of BPH does not increase the risk of prostate cancer when age, race, and body mass index were all taken into consideration.<sup>[8]</sup>

### *Genetic associations between BPH and PCa*

Extensive genetic studies into the pathophysiology of prostate cancer have been conducted with the realistic potential to provide bespoke care for this group of patients depending on the genetic and epigenetic anomalies they harbor<sup>[9,10]</sup> [Table 2]. It has been over a decade since genetic overlapping between these two diseases was identified.<sup>[11]</sup> Prakash *et al.*<sup>[11]</sup> with the use of 511 non-redundant genes, identified clusters of proliferation genes to be associated with BPH and co-existing oncogenes identified in both BPH and PCa tissues. However, the genetic and epigenetic insight into BPH has somewhat lagged behind relatively in comparison to the progress that has been made with PCa.<sup>[11]</sup>

With this in mind, recent studies have further re-iterated the link between BPH and prostate cancer with genetic evidence. With the utilization of technology, such as next-generation sequencing, it has provided us the ability to uncover disease-related variants by processing large genomic intervals in a rapid and thorough manner. Mutations in the HBOX13 gene have been established to be associated with the development of prostate cancer. Saaristo *et al.*<sup>[12]</sup>

**Table 1: A summary of contemporary landmark epidemiological studies assessing the risk of PCa in the presence of a diagnosis of BPH**

Group	Published	BPH patient no.	Period	Summary of outcome
Hung <i>et al.</i> <sup>[7]</sup>	2013	4736	1997-2008	Higher OR of 26.2 of developing PCa
Orsted <i>et al.</i> <sup>[6]</sup>	2011	3,009,258	1980-2007	2-3 and 2-8 times increased risk of PCa incidence and mortality, respectively
Schenk <i>et al.</i> <sup>[8]</sup>	2011	1,549	1993-2003	No association
Chokkalingam <i>et al.</i> <sup>[4]</sup>	2003	86, 626	1964-1983	1.2 and 1.7 times increased risk of PCa incidence and mortality, respectively

BPH=Benign prostatic hyperplasia, OR=Odds ratio, PCa=Prostate cancer

**Table 2: Common genetic anomalies shared between BPH and PCa**

Group	Gene	Function
Klotsman <i>et al.</i> <sup>[14]</sup>	SRDAR2	Implicated in androgen conversion
Tang <i>et al.</i> <sup>[15]</sup>	CYP17	Implicated in androgen synthesis
Habuchi <i>et al.</i> <sup>[16]</sup>	CYP19	Implicated in androgen degradation
Henrique <i>et al.</i> <sup>[18]</sup>	14-3-3 $\sigma$	Tumor suppressor involved in cell cycle regulation and apoptosis
Bastian <i>et al.</i> <sup>[19]</sup>	RASSF1A	Tumor suppressor gene
Henrique <i>et al.</i> <sup>[20]</sup>	MT1G	Implicated in heavy metal binding
Enokida <i>et al.</i> <sup>[21]</sup>	MDR1	ABC transporter which transports various molecules across cell membranes

BPH=Benign prostatic hyperplasia, PCa=Prostate cancer, SRDAR2=Steroid 5- $\alpha$  reductase type II, CYP=Cytochrome P450, RASSF1A=Ras association domain-containing protein 1, MT1G=metallothionein 1G, MDR1=Multi-Drug resistant protein 1

assessed 100 single-nucleotide polymorphisms (SNP) in over 500 men with BPH who either remained with this disease or later developed prostate cancer. As expected, a mutation in HBOX13 was strongly associated with the prostate cancer specimens. However, BPH patients carrying this mutation had a 4.6 times greater risk of developing prostate cancer compared to non-carriers (OR 4.56, CI 95% 1.29-16.11,  $P = 0.0098$ ). Interestingly, the average PSA levels at baseline biopsy in both groups were very similar (BPH = 7.3  $\mu\text{g/l}$  (range 0.5-44  $\mu\text{g/l}$ ) and 8.0  $\mu\text{g/l}$  (range 2.1-75  $\mu\text{g/l}$ ) respectively). This is the first study to report genetic predisposition to developing PCa in BPH patients and gives a future potential of predicting those men with BPH who will later go on develop PCa.

Similarly, variants of numerous other genes have been shown to confer an increased risk of the development of both BPH and prostate cancer.<sup>[13]</sup> Variants of genes implicated in androgen conversion within the prostate (SRDAR2), those involving androgen synthesis (CYP17) and degradation (CYP19) have all commonly been shown to increase the risk of the development of BPH and prostate cancer.<sup>[14-16]</sup> DNA methylation is an epigenetic method, in which genes can be regulated via silencing of that particular portion of the genome. In PCa, there are numerous genes that have been shown to be hypermethylated and hence inactivated, ultimately leading to the progression of prostate cancer.

14-3-3 $\sigma$  is a putative tumor suppressor gene involved in cell cycle regulation and apoptosis following DNA damage. Its loss of expression has been established in numerous cancers including PCa.<sup>[17]</sup> With the use of 189 patient specimens, Henrique *et al.*<sup>[18]</sup> have shown that via epigenetic mechanisms, 14-3-3 $\sigma$  is significantly hypermethylated in both PCa and BPH samples. Similarly, epigenetic phenomenon have been shown to be shared between BPH and PCa tissues including tumor suppressors genes (RASSF1A),<sup>[19]</sup> heavy metal binding genes (MT1G),<sup>[20]</sup> or genes which encode for proteins such as ABC transporters (MDR1).<sup>[21]</sup> All of these genes have also been shown to be hypermethylated in BPH and PCa specimens, suggesting the existence of an epigenetic link between BPH and PCa.

Despite the progress made in the genetic and epigenetic insight in to prostatic disease, to-date, there are no genetic markers that are currently utilized in routine general clinical practice for either BPH or prostate cancer.<sup>[13]</sup>

### Inflammatory associations of BPH and PCa

Inflammation has been detrimentally linked to the development and presence of both BPH and PCa. Large population-based trials such as the Medical Therapy of Prostate Symptoms (MTOP) and more recently the Reduction by Dutasteride of Prostate Cancer Events (REDUCE) studies have demonstrated the association between inflammation and development of BPH.<sup>[22,23]</sup> The MTOP study demonstrated that approximately 40% of baseline histological specimens of BPH harbored chronic inflammatory infiltrates, and this group of men had an increased baseline prostatic size (41 mls vs. 37 ml), higher baseline PSA levels (3.1 ng/ml vs. 2.3 ml), greater chance of disease progression (21% vs. 13.2%), increase in IPSS score (13.7% vs. 11.2%), and a significant increase in the incidence of the development of acute urinary retention (5.6% vs. 0%) in comparison to those men who harbored no stigmata of chronic inflammation on baseline histological specimens. The REDUCE trial showed that chronic inflammatory infiltration was more prevalent in men with BPH in comparison to benign controls and furthermore, the degree of LUTS attributed to BPH was correlated with the severity of inflammation present.

With the use of immune-histochemical analysis in 282 specimens collected from a more focused group of BPH sufferers, all of whom underwent surgical treatment

of the disease, demonstrated that 82% and 81% of specimens had T-lymphocyte markers and macrophage markers, respectively.<sup>[24]</sup> Within this cohort, a correlation existed between those with high-grade inflammation and a higher IPSS score (21 vs. 12;  $P = 0.02$ ) and prostatic volume (77 cm (3) vs. 62 cm (3);  $P = 0.002$ ).

Chronic inflammation is thought to be one of the numerous driving factors for the development of PCa. McLennan *et al.*<sup>[25]</sup> performed needle biopsy of the prostate at baseline and after a 5-year period and reported that the 5-year incidence of prostate cancer was 20% in men harboring inflammation at baseline compared to 6% in those with no inflammation at baseline biopsy. Genes associated with inflammation have also demonstrated to be aberrantly expressed resulting in a pro-tumorigenic outcome in PCa. GSTpi are thought to have an important role in cancer prevention by providing cellular protection and subsequent inflammatory reaction. This is achieved by preventing cellular damage from free radicals and other carcinogenic compounds. This gene has been shown to be down-regulated in PCa and the loss of which may be responsible in patients with genetic predisposition for the transition of PIN into high-grade intraepithelial neoplasia (HGPIN) and PCa.<sup>[26,27]</sup> Similarly, SNPs within the MSR1 genes, which encodes for macrophage scavenger receptor 1, have been shown to be significantly associated with prostate cancer.<sup>[28]</sup>

Despite compelling evidence that both BPH and PCa share common inflammatory component driving disease progression, there is very limited evidence that the use of non-steroidal anti-inflammatory (NSAIDs) agents has a favorable outcome in decreasing incidence of BPH. A recently reported study investigated the use of NSAIDs in the form of aspirin and ibuprofen in nearly 5000 men.<sup>[29]</sup> It concluded that the use of these agents had no protective effect on the development of BPH and LUTS.

There is limited evidence that the use NSAIDs may provide a protective effect in the development of prostate cancer. A recent population study using over 9000 men has reported a modest reduction of 10% in the development of PCa with men using regular NSAID agents.<sup>[30]</sup> The same group of authors also published a meta-analysis on this topic analyzing over 24,000 patients.<sup>[31]</sup> The outcome of the meta-analysis was that the use of NSAID agents with the view of preventing prostate cancer was a suggestive one but certainly not one that is conclusive and would require further well-designed observational studies for greater insight and understanding into this topic.

#### *Hormonal associations between PCa and BPH*

A common and well established driving factor co-linking the advancement and progression of both BPH and PCa is the action androgens have on prostatic tissue. Not only are androgens explicitly required for pathological

growth of the prostate, they are vital for normal growth and development of this gland. The iso-enzymes 5- $\alpha$  reductase type 1 and 2 convert testosterone into its active form dihydrotestosterone (DHT), which in turn facilitates cellular differentiation and proliferation of prostatic tissue.<sup>[32]</sup> Abnormal prostatic growth occurs when there is a disturbance on the androgen profile of the prostate, which encourages a proliferative state of the gland. This disrupted androgen cycle overcomes apoptosis within the prostate, therefore, disrupting the homeostatic regulation of prostate cell proliferation and cell death. The suppression of androgen activity on the prostate forms the basis of pharmacological treatment of both BPH and PCa.

The hypothesis that higher circulating androgens may confer increased risk of prostate cancer has been assessed by large trials investigating the action anti-androgen agents have on the development of PCa.<sup>[33,34]</sup> The Prostate Cancer Prevention Trial concluded that prescribing men finasteride for 7 years resulted in reducing the risk of the development of PCa by 25%.<sup>[34]</sup> Similarly, the REDUCE trials, which prescribed men dutasteride over a 4-year period, resulted in a decrease in the incidence of PCa by 23%.<sup>[33]</sup> As expected, both studies reported a favorable outcome on lower urinary tract symptoms related to BPH.

## CONCLUSION

Genetic, hormonal, and inflammatory mechanisms have all been shown to be common pathophysiological driving mechanisms for the development of both BPH and PCa, thus linking these diseases together. However, on a cellular and molecular level, no study has shown that the development of BPH tissue has later converted into an oncological disease. Furthermore, the exact pathways of these prostatic diseases have yet to be fully understood. This is essential for us to optimize future treatment strategies for both diseases and an issue that future basic science studies should target.

Epidemiological data suggests that the presence of BPH increases the risk for a man to develop PCa in his lifetime. However, a causal link cannot be established from the current data. To determine the risk of the development of PCa of men presenting with BPH, we require further detailed epidemiological studies, which would ideally include a large cohort of men from a global platform.

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