


Research Paper

A seemingly most effective target for early detection and intervention of prostate tumor invasion

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Abstract

This commentary proposes that budding tumor cell projections from focally disrupted tumor capsules represent a most effective target for early detection and intervention of prostate tumor invasion. The rationale, supporting data, and clinical applications of the hypothesis are discussed.

Key words: Prostate cancer; Invasion; Early detection; Basal cells; Tumor cell budding; Stem cells

1. The structural features of the human prostate

The normal and pre-invasive prostate tumor epithelium, which is the histological origin of over 80% of the human prostate malignancies, is physically separated from the stroma by the basal cells and basement membrane. Basal cells are joined by intercellular junctions and cell surface adhesion molecules, forming a largely continuous sheet encircling epithelial cells [1-2]. The basement membrane is composed of type IV collagen, laminins, and other molecules, forming a continuous lining surrounding the basal cell layer [3-4] (Fig 1). Together, the basal cell layer and the basement membrane constitute a morphologically distinct capsule, which is a permanent structural element largely independent of hormonal regulation and the host's bio-physiological condition [1-4]. The epithelial cells are held in place by intercellular junctions and cell surface adhesion molecules (Fig 2). The epithelium is normally devoid of blood vessels and lymphatic ducts and totally relies on the stroma for its basic needs. Due to these structural relationships, the physical and functional status of the tumor capsule significantly impacts the biological

behavior of the epithelial cells, and the disruption of the tumor capsule is a pre-requisite for prostate tumor invasion and metastasis.

2. Focal basal cell alterations and disruptions in tumor capsules

Prostate tumor invasion is believed to be a multistage process, progressing sequentially from normal to hyperplasia, to prostatic intraepithelial neoplasia (PIN), and to invasion [5-8]. Progression from PIN to invasion is believed to be triggered by overproduction of proteolytic enzymes primarily by cancer cells, which cause degradation of the tumor capsule [9-10]. These theories are consistent with experimental data from tissue cultures or animal models, but they are hard to reconcile with four critical facts: (a) previous studies have revealed that some healthy men between 19 and 29 years old demonstrated a spectrum of proliferative abnormalities in the prostate, including atypical hyperplasia, dysplasia, and incipient adenocarcinoma [11-13], (b) recent studies have detected a DNA phenotype that is identical to the DNA structure of invasive prostate cancer in certain healthy men, and in normal prostate tissues adjacent to prostate cancer [14-17], (c) a majority of PIN express high le-

vels of proteolytic enzymes, but only about 30% of untreated PIN progress to invasive lesions during patients' lifetime [18-21], and (d) results from worldwide human clinical trials with a wide variety of proteolytic enzyme specific inhibitors to treat or prevent tumor invasion have been very disappointing [22,23]. Together, these suggest that alternative mechanism(s) may exist for prostate tumor invasion.

A number of clinic tests, including the measurement of the serum level of prostate specific antigen (PSA), MRI, and gene expression profiling, are increasingly used in the clinic, whereas none of those could predict which of the PIN lesions will progress [24-29]. The only established approach to monitor PIN progression is repeat biopsy [25-29], which is costly and painful. Since over 90% of prostate cancer related mortality result from invasion-related illnesses, and the incidence of PIN could be up to 16.5%-25% in routine or ultrasound guided prostate biopsies [25-29], there is an urgent need to uncover the intrinsic mechanism of tumor invasion, and to develop novel approaches to identify the specific precursor of invasive prostate lesions.

Promoted by the fact that the basal cell layer is the sole source of tumor suppressor p63 and maspin in prostate [31-33], and that the absence of the basal cell layers is one of the most distinct morphological signs of invasive cancers, our recent studies have attempted to identify the early signs of tumor capsule disruptions. Our initial study examined the physical integrity of basal cell layers in 50 patients with co-existing pre-invasive and invasive prostate tumors. Of 2,047 ducts and acini examined, 197 were found to harbor focal disruptions (the absence of basal cells resulting in a gap greater than the combined size of at least 3 basal cells) in the tumor capsule. The frequency of focal tumor capsule disruptions varied from none in 22 (44%) cases to over 1/3 of the ducts or acini with focal disruptions in 17 (34%) cases [34-38].

Compared to their non-disrupted counterparts, focally disrupted tumor capsules displayed a number of unique alterations, including a significantly lower proliferation index and p63 expression, but a significantly higher frequency of degeneration, apoptosis and infiltration of leukocytes, which are generally located at or near focally disrupted tumor capsules (Fig 3d).

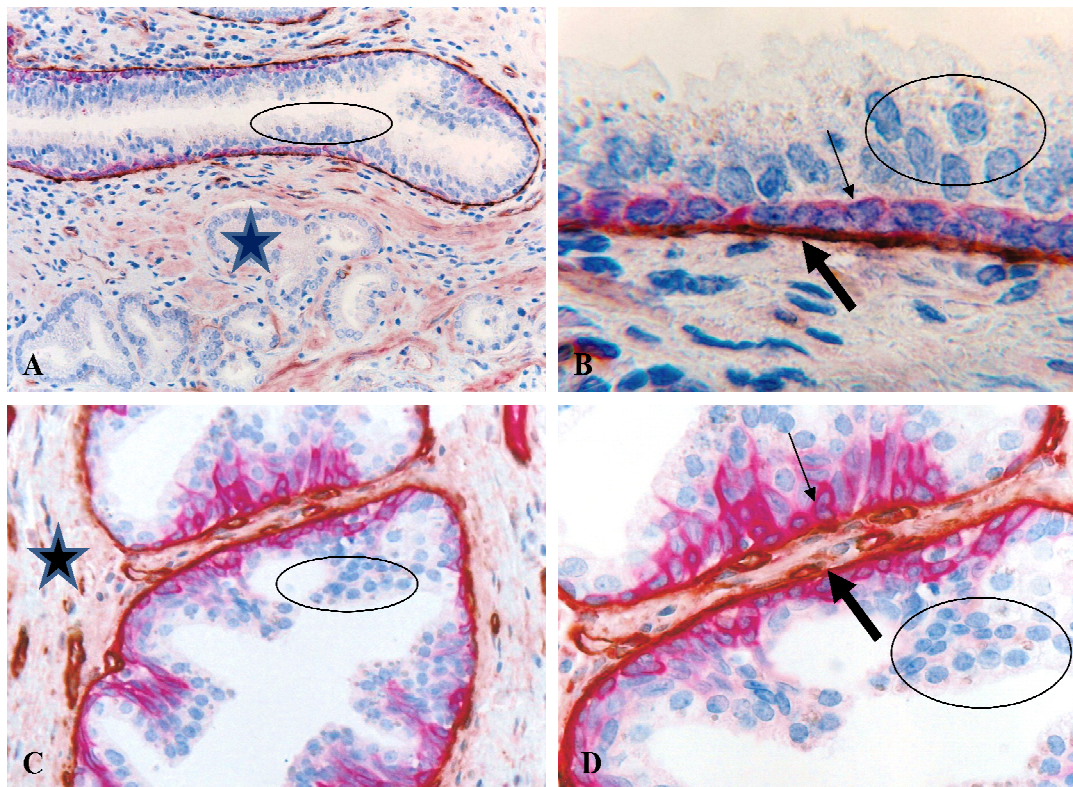


Fig 1. Structural relationships among the prostate epithelium, basal cell layer, basement membrane, and stroma. Human prostate sections were double immunostained for cytokeratin 34BE12 (red) and collagen IV (brown). Circles identify epithelial cells. Stars identify the stroma. Thin arrows identify the basal cell layer. Thick arrows identify the basement membrane. A and C: 150X. B and D: a higher magnification (400X) of A and C, respectively.

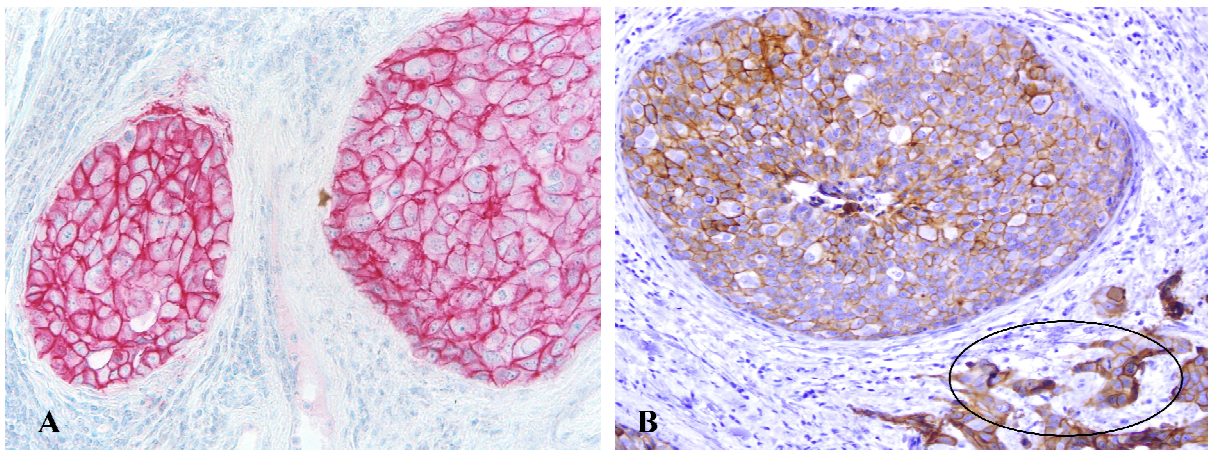


Fig 2. E-cadherin expression in epithelial cells. Sections were immunostained for cell surface adhesion molecule E-cadherin. Note that both the pre-invasive and invasive cancer cells (circle) show strong E-cadherin expression, and the edges of pre-invasive tumors are smooth without budding cells. 150X.

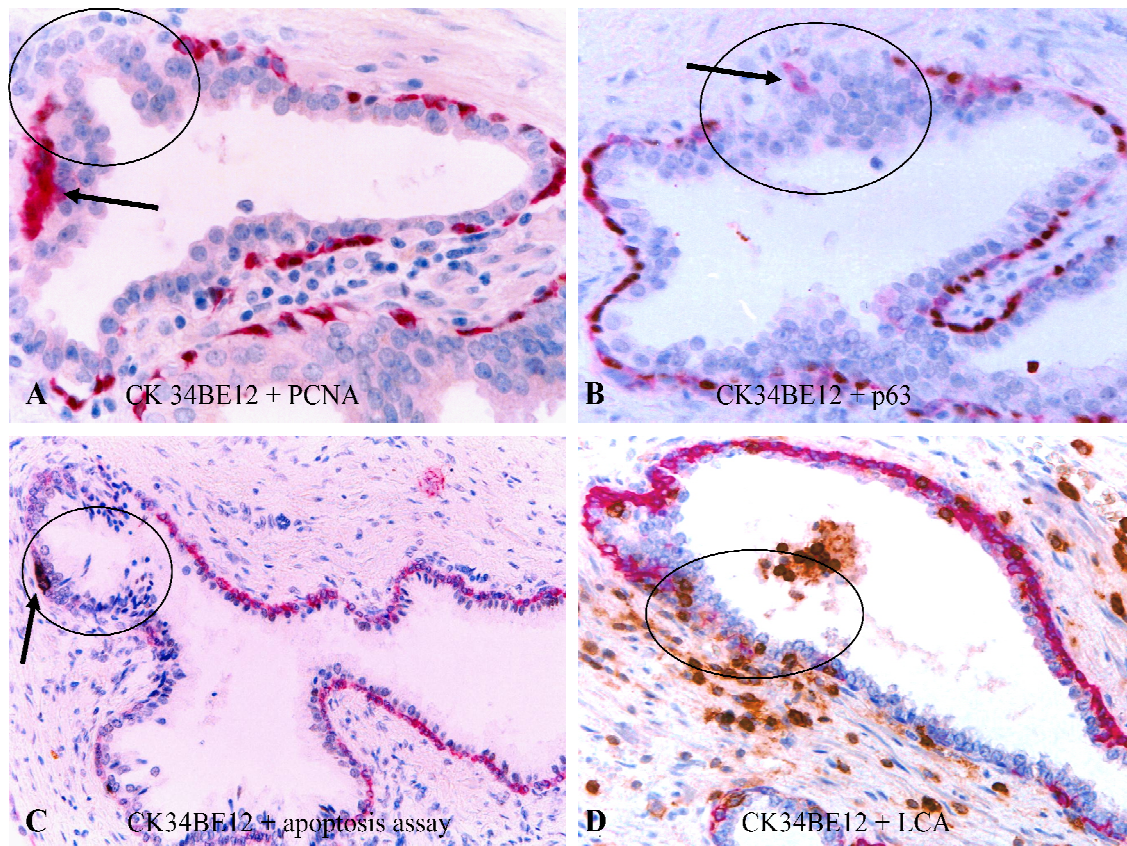


Fig 3. Focal disruptions in tumor capsules. Human prostate sections were double immunostained with different markers. Circles identify focal disruptions in the tumor capsules. Arrows identify degenerated (A), isolated (B), and apoptotic (C) basal cells. Note that most leukocytes (brown particle-like structures) are located at or near focally disrupted capsules. 200X.

3. Tumor cell budding from focally disrupted tumor capsules

Focal disruptions in the tumor capsule appear to significantly impact the morphological and biological presentations of associated epithelial cells. Although

most focal tumor capsule disruptions were seen in PIN, a subset of normal- and hyperplastic-appearing ducts and acini also showed focal disruptions in their capsules. As shown in Fig 4, epithelial cells overlying focally disrupted capsules of two morphologically normal appearing ducts are arranged as finger- or

tongue-like projections budding from focally disrupted capsules. Budding cell projections often punctured deep into the stroma and were morphologically indistinguishable from invasive cancer cells (Fig 4). A majority of epithelial structures with budding cell projections were adjacent to invasive lesions and all budding cells from the same or different cases shared a similar morphological and immunohisto-

chemical profile with a substantially higher proliferation index than their parent structures [34-38].

These budding cell projections also expressed high levels of malignancy-associated markers alpha-methylacyl-CoA racemase (AMACR) and prostate specific antigen (PSA). In contrast, cells within the same duct or acinus but distant focally disrupted tumor capsules were largely devoid of these two molecules (Fig 5).

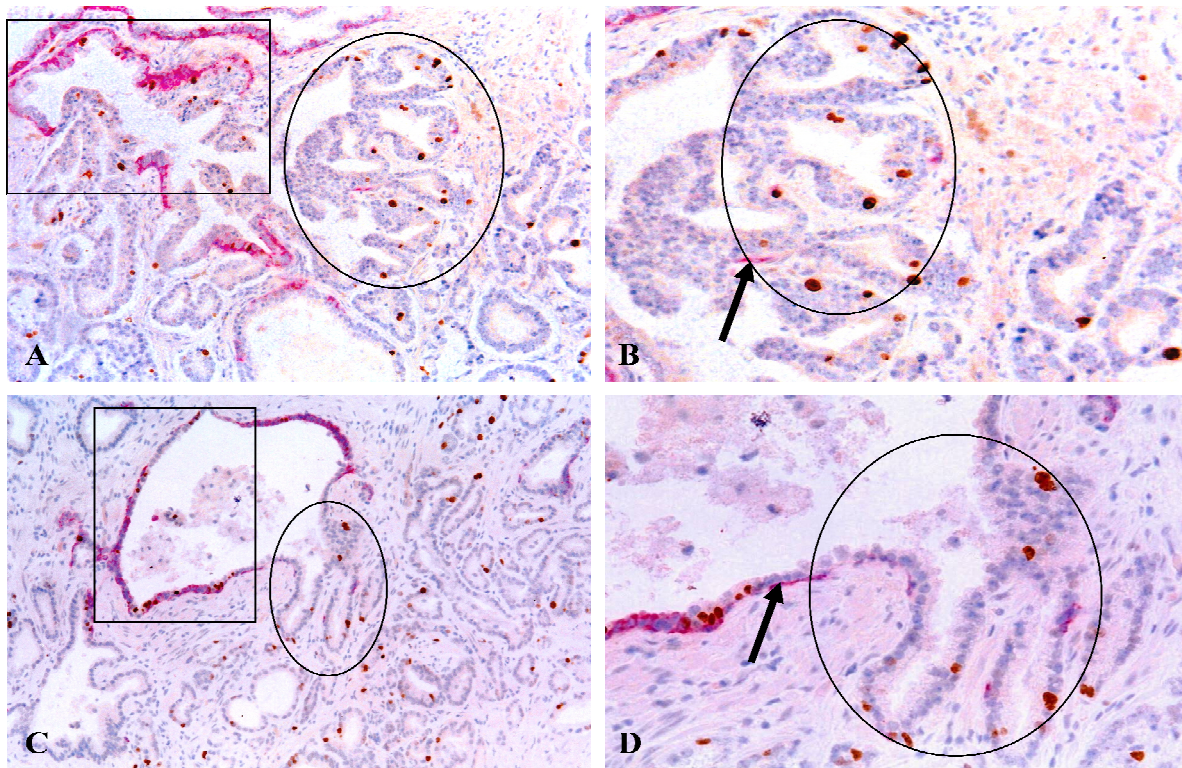


Fig 4. Tumor cell budding from focally disrupted tumor capsules. Human prostate tumor tissue sections were double immunostained for CK34BE12 (red) and Ki-67 (brown). Circles identify budding tumor cell projections. Squares identify normal appearing prostate ducts in which the budding cell projections are originated. Arrows identify basal cells. A&C: 100X. B&D: 300X.

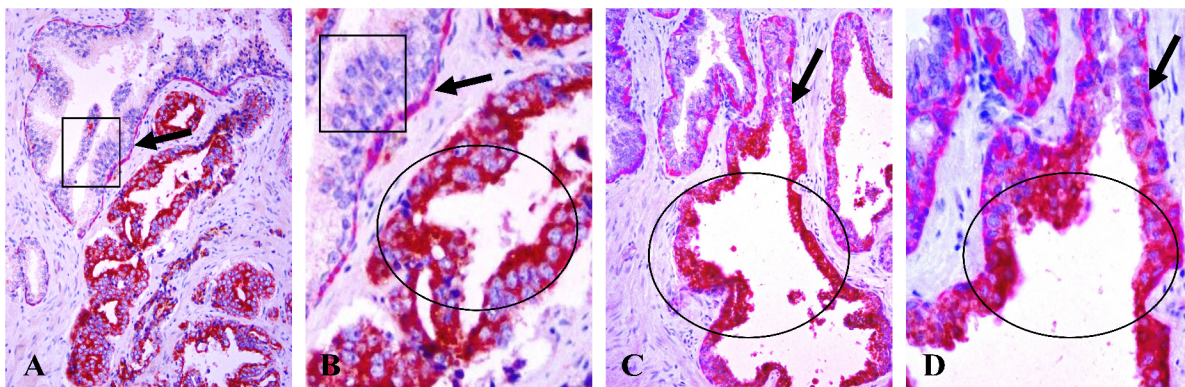
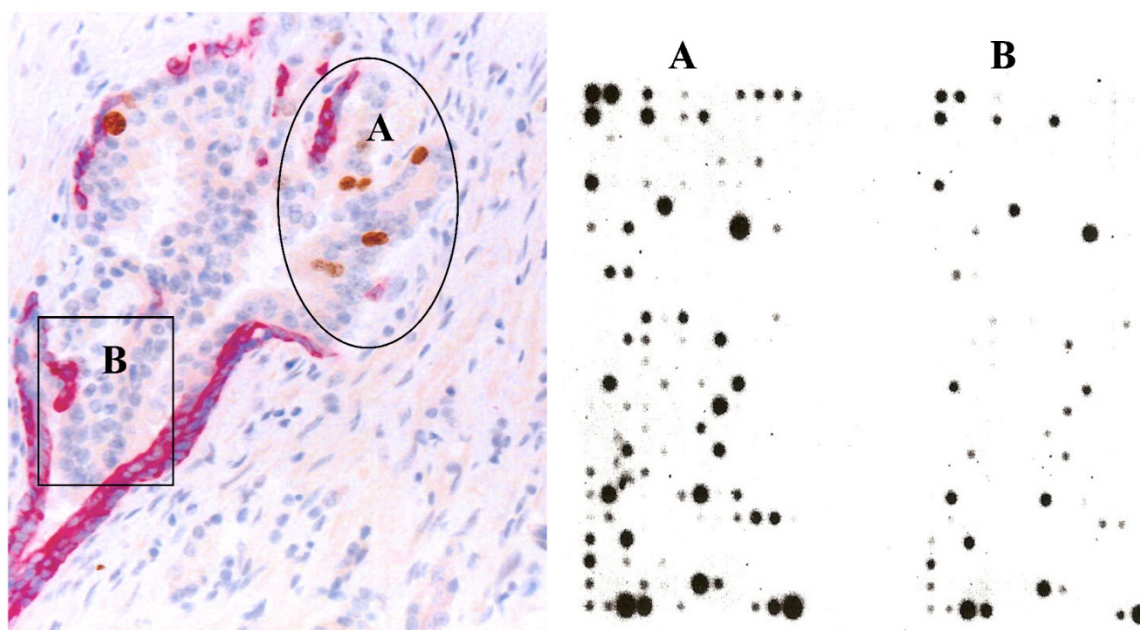


Fig 5. Expression of malignancy-associated molecules. Human prostate tumor sections were double immunostained for CK34BE12 (red) and AMACR (A-B; brown) or PSA (C-D; brown). Circles identify budding cell projections with AMACR or PSA expression. Squares identify morphologically similar cells without the expression of AMACR or PSA. Arrows identify residual basal cell layer. A & C: 100X. B & D: a higher magnification (300X) of A&C, respectively.



	Gene name	Functional grouping	Fold change
1	LIF	Growth factors	47.37
2	MCL1	Anti-apoptosis	6.72
3	TNFRSF7	Anti-apoptosis	7.91
4	KIT	Stem cell lineage marker	5.03
5	NCOR2	Stem cell-related marker	5.45
6	ENG	Endothelial cell marker	6.38
7	ICAM2	Endothelial cell marker	12.12
8	KRT17	Epithelial cell marker	7.15
9	ITGA3	Cell-matrix adhesion	5.52
10	ITGB3	Cell-matrix adhesion	7.14
11	CCL2	Chemokine, cytokine, & receptor	14.33
12	CX3CL1	Chemokine, cytokine, & receptor	6.14
13	CCR1	Chemokine, cytokine, & receptor	5.19
14	CXCR4	Chemokine, cytokine, & receptor	12.81
15	TNFRSF10D	TNF receptor family	8.20
16	TNFRSF12A	TNF receptor family	5.35
17	TNFRSF25	TNF receptor family	8.52
18	TIMP1	ECM inhibitor	5.25
19	TIMP3	ECM inhibitor	7.87
20	MMP26	Extracellular matrix proteins	-6.94
21	IL10	Interleukin & receptor	-9.50
22	IL12RB2	Interleukin & receptor	-7.02
23	IL6R	Interleukin & receptor	-7.24

Fig 6. Differential gene expression between cells overlying focally disrupted tumor capsule (A) and cells within the tumor core (B). Note that budding cells from focally disrupted tumor capsule have a significantly higher expression of invasion and stem cell related genes.

Gene expression profiling with RNA extracted from microdissected budding cell projections and their adjacent counterparts within the same tumor showed that budding cell projections had a significantly higher expression of stem cell- and tumor invasion-related genes [36] (Fig 6).

4. Clinical implications of tumor cell budding from focally disrupted tumor capsules

Our findings of tumor cell budding from focally disrupted prostate tumor capsule are consistent with those of a number of previous studies in human esophageal and colorectal cancers, which detected a similar pattern and frequency of tumor cell budding, and revealed that tumors with budding cells were significantly correlated with invasion, metastasis, and worse prognosis [39-41]. The results of our gene expression profiling are also in line with those of a recent study, which showed that microdissected cells from the periphery and the center of the same *ductal carcinoma in situ* had a markedly different frequency and pattern in the expression of 22 genes, assessed with Atlas human Cancer 1.2 Arrays containing 1176 known genes [42].

Together, our findings suggest that tumor cell budding from focally disrupted tumor capsules is likely to represent an early sign of tumor invasion. Our findings also suggest that budding tumor cell projections are likely to result from monoclonal proliferation of an activated tumor progenitor or a biologically more aggressive cell clone and represent the direct precursors of invasive prostate lesions. If further validated, budding tumor cell projections from focally disrupted tumor capsules could be considered as a most effective target for early detection and intervention of prostate tumor invasion for three main reasons. **First**, as focal capsule disruptions with budding cells are also seen in normal- or hyperplastic-appearing ducts and acini, budding cell projections may also represent an early sign of prostate tumor progression that eventually leads to invasion. Thus, the development of more feasible technical approaches to detect focal capsule disruptions with budding epithelial cells in biopsy samples may significantly facilitate differentiation between clinically aggressive and indolent pre-invasive prostate lesions, and also leads to identification of the specific individuals at increased risk to develop invasive prostate cancer. **Second**, as these budding cell projections are generally surrounded by or adjacent to leukocytes and other stromal cells (as shown in Fig 3), the degradation products from the tumor capsules and the secretory products from the budding cells are likely to enter the blood stream. Thus, the development of a

quantitative assay to measure the levels of these molecules in the blood samples could be potentially used as a screen tool to identify the individuals at greater risk to develop invasive prostate cancer. **Third**, as all budding cell projections within the same or different cases are morphologically and immunohistochemically similar, they are likely to have a very similar molecular profile and also share the same pathway for invasion. Thus, the identification of the shared molecules within these cells could lead to the development of effective therapeutic agents to specifically target all these cells, which could potentially block tumor invasion at the very early stage. However, it is currently difficult, if not impossible, to fully elucidate the molecular and biochemical profiles of these very limited budding cells, due to the lack of sensitive technical approaches.

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Conflict of Interest

The author has declared that no conflict of interest exists.

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