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Prominence of urinary biomarkers for bladder cancer in the COVID-19 era: From the commercially available to new prospective candidates

Xuan-Mei Piao¹⁽¹⁾, Howon Kang^{1,2}⁽¹⁾, Wun-Jae Kim^{1,3}⁽¹⁾, Seok Joong Yun^{1,2}⁽¹⁾

¹Department of Urology, College of Medicine, Chungbuk National University, Cheongju, ²Department of Urology, Chungbuk National University Hospital, Cheongju, ³Institute of Urotech, Cheongju, Korea

Molecular markers detected in urine may improve our understanding of the evolution of bladder cancer (BCa) and its micro- and macroenvironment. Detection of such markers will identify disease earlier, allow stratification of patients according to risk, and improve prognostication and prediction of outcomes, thereby facilitating targeted therapy. However, current guidelines have yet to embrace such markers for routine management of BCa, and most research studies have focused on urine-based tumor markers. In this review, we summarize known urinary biomarkers for BCa and highlight newly identified molecules. We then discuss the challenges that must be overcome to incorporate these markers into clinical care.

Keywords: Biomarkers; DNA; Protein; RNA; Urinary bladder neoplasms

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INTRODUCTION

Bladder cancer (BCa) is a common disease worldwide, with high morbidity [1,2] however, developments in medical technology mean that it is no longer fatal in many cases. Despite this, mortality rates for those with muscle-invasive bladder cancer (MIBC) or those who do not receive optimal treatment remain high [3]. One of the hallmarks of BCa is its heterogeneity, which makes it difficult to manage [4]. This heterogeneity is due to genetic, transcriptomic, epigenetic, and/or phenotypic changes, which result in a molecularly heterogeneous tumor comprising cancer cells with diverse molecular signatures [4,5]. Therefore, tumor heterogeneity is the major barrier to successful management of BCa, including early identification of non-muscle-invasive bladder cancer (NMIBC) or MIBC. Appropriate and early identification will enable suitable treatment planning and assessment of prognosis. Consequently, identification of suitable and reliable tumor biomarkers is essential for diagnosis, prognosis, and treatment planning [6].

Diagnosis and follow-up of BCa are dependent on cystoscopy. This is a highly invasive procedure, which itself can cause complications such as infection or hematuria. Moreover, the high recurrence rate of BCa, along with the frequent requirement for surveillance, place huge economic and quality-of-life burdens on patients [7]. Voided urine cytology (VUC), which has been applied to BCa diagnosis, is a noninvasive option; however, its use is restricted by relatively low

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Department of Urology, Chungbuk National University Hospital, College of Medicine, Chungbuk National University, 776, 1sunhwan-ro, Seowon-gu, Cheongju 28644, Korea

TEL: +82-43-269-6371, FAX: +82-43-269-6144, E-mail: sjyun@chungbuk.ac.kr

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sensitivity (particularly for diagnosing low-grade tumors); indeed, a previous study reported sensitivity of only 25% [8]. Thus, efforts have been made to explore cost-effective and noninvasive alternatives to cystoscopy. In particular, delivering cancer care during the present COVID-19 pandemic is challenging given the competing risks of cancer-specific death vs. a potentially lethal coronavirus infection. This highlights an urgent need to develop a guide to pragmatic management of BCa.

Clinical decision-making in a pandemic requires a balance between the probable benefits and risks; patients should attend hospitals only when strictly necessary, but care must be given to those most in need. Given this, the European Association of Urology developed guidelines suggesting a traffic-light surveillance pathway based on primary tumor grade and the presence of hematuria [9,10]. This guideline recommends that patients with low-risk or intermediate-risk tumors, and who are asymptomatic, wait a further 6 months for cystoscopy [11] Although the guidelines are adaptable to the current situation, some patients are happy to defer this often costly and painful process; however, others would rather undergo the procedure quickly than worry about the ambiguity of their disease status.

This period of uncertainty requires timely action and innovation. Urinary biomarkers would enable early detection of cancer, particularly in patients for whom cystoscopy has been deferred in accordance with the up-to-date guidelines. In this review, we summarize the urinary biomarkers used for BCa diagnosis and discuss research and development of new advanced biomarkers.

SIGNIFICANCE OF BIOMARKER RESEARCH

1. Bladder cancer markers: what for?

Tumor markers are molecules secreted directly by tumor cells or indirectly by other cells in response to a tumor [12]. Biomarkers can be used for screening, diagnosis, monitoring/surveillance, and prognosis. However, there are no currently accepted biomarkers for BCa screening; therefore, the gold standard tests (a combination of VUC and cystoscopy) are still used for diagnosis in practice [13]. The main role of surveillance markers is to reduce the need for invasive cystoscopy; however, like diagnostic markers, they are not sufficiently reliable for routine clinical use [13,14]. Prognostic markers can be used to stratify patients according to clinical outcome (e.g., recurrence or progression), thereby helping clinicians decide which treatments are most beneficial in a particular case [15]. Consequently, the main goal for those

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developing biomarkers is to identify relevant molecules or tests that can improve clinical decision-making in a costeffective way.

2. Biomarker sources: liquid biopsy vs. tissue biopsy

Tissue biopsy is the traditional approach used for cancer diagnosis. The analysis of biopsy samples detects abnormal tumor cells in tumor-like tissue and surrounding tissue. However, this procedure is highly invasive, painful. expensive, and time-consuming. In addition, it requires the intervention of a skilled clinician owing to difficulties in obtaining the right sample for analysis [16]. In addition, the heterogeneity of BCa is a critical limitation in that a tissue biopsy may not always reflect the entire tumor landscape. The tumors within a bladder vary with respect to morphology, gene expression profile, and mutations. This heterogeneity takes several forms: (1) patient-to-patient (interpatient) heterogeneity; (2) spatial region-to-region variation within a tumor in the same patient (intra-tumoral heterogeneity); and (3) tumor-to-tumor variation, including primary tumor to primary tumor, primary tumor to metastatic site, and metastatic site to metastatic site in the same patient (intertumoral heterogeneity) [4]. Consequently, tissue obtained from different areas of the same tumor, from different sites (primary/metastatic) within the same patient, or from different patients may harbor radically different mutations and gene expression patterns [17]. Thus, the use of minimally invasive procedures such as liquid biopsies is gaining traction [18]. Liquid biopsies, which work by measuring circulating tumor-derived material such as circulating tumor cells (CTCs), circulating cell-free tumor DNA (ctDNA), circulating cell-free tumor RNA (ctRNA), proteins, and extracellular vesicles (EVs) in body fluids, have great potential to overcome the limitations inherent to tissue sampling [18]. Because liquid biopsies are minimally invasive, the risk of complications and pain is reduced. Most importantly, liquid biopsies may better represent tumor heterogeneity and allow monitoring of changes in real time. Although liquid biopsies require minimal medical skills and surgical facilities, they still require specialized laboratory equipment and qualified personnel. Another limitation of liquid-biopsy-based research is that it lacks standard protocols, and the low concentrations of materials may complicate interpretation of the results [19]. Nevertheless, the ease of sample collection and the possibility of time-independent analysis are tremendous assets. Ideally, use of reliable urinary biomarkers for BCa will facilitate patient management and could even provide an at-home service. Tissue and liquid biopsies are depicted and

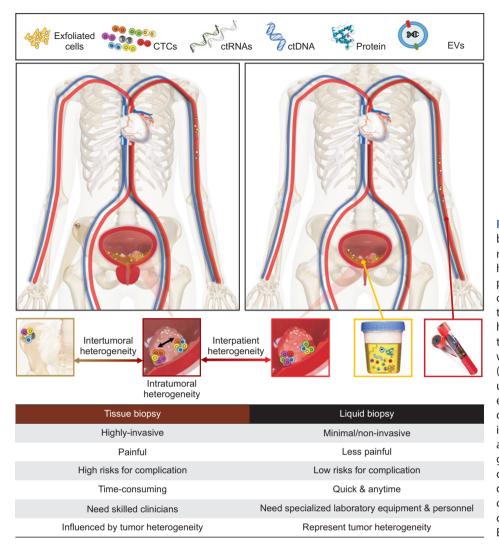


Fig. 1. Comparison of tissue and liquid biopsies. Heterogeneity of BCa exists not only between patients (interpatient heterogeneity) but also within the same patient. Intratumoral heterogeneity is caused by variations within regions of the same tumor, whereas intertumoral heterogeneity refers to differences between multiple tumors or metastases within a single patient. Liquid biopsies (which comprise mainly blood and urine) may better represent these heterogeneities. Analysis of liquid biopsies detects alterations in levels of circulating tumor proteins, ctDNA, ctRNA, CTCs, and tumor/normal cell-derived EVs, giving them several advantages over conventional tissue biopsies. BCa, bladder cancer; CTCs, circulating tumor cells; ctDNA, circulating cell-free tumor DNA; ctRNA, circulating cell-free tumor RNA; EVs, extracellular vesicles.

compared in Fig. 1.

URINARY BIOMARKERS FOR BLADDER CANCER

1. Urine as a source of biomarkers

The majority of tumor markers are secreted into blood and can be measured in blood; however, they can also be measured in other types of liquid biopsy (e.g., saliva, urine or seminal plasma, and tissues). Some markers are specific to a single type of cancer, whereas others are associated with several types of cancer [12,20]. Unfortunately, unlike the prostate, which secretes prostate-specific antigen into serum, the bladder secretes no organ-specific markers. The urinary bladder is a hollow muscular organ that stores urine (the capacity is about 300–500 mL). It is a small organ, and most BCa tumors are usually less than several centimeters in size; thus, markers secreted by the tumor may not be easy to detect in blood. However, because bladder tumors are in direct contact with urine, many studies focus on identification of urinary biomarkers of BCa. To date, six urine-based methods have been approved by the U.S. Food and Drug Administration (FDA) for clinical use; however, they must still be used in combination with cystoscopy, and their use remains controversial [21,22]. These FDA-approved markers/tests are the Nuclear matrix protein 22 (NMP22) quantitative and qualitative tests, the bladder tumor antigen (BTA) STAT/ TRAK (Polymedco, New York, NY, USA), the ImmunoCyt/ uCyt+ assay (DiagnoCure Inc, Quebec, QC, Canada), and the UroVysion bladder cancer kit (Abbott Molecular Inc, Chicago, IL, USA). Most studies of urine-based biomarkers for BCa are of emerging biomarkers to be applied in the future.

2. FDA-approved urinary biomarkers for bladder cancer

NMP is a nonchromatin structure responsible for regulating DNA replication, transcription, and RNA processing [23-25]. Expression of NMP22 is increased in urothelial tu-

mors, and shedding of apoptotic tumor cells into the urine enables detection of this protein in body fluid. However, NMP22 is also present in normal urothelial cells. NMP22 tests include a quantitative enzyme-linked immunosorbent assay (ELISA; the NMP22 test; Abbott) and a qualitative point-of-care (POC) test (NMP22 BladderChek test; Abbott), which are designed to detect the NMP22 antigen in urine, thereby assisting both BCa diagnosis and monitoring of BCa recurrence. The tests are painless and noninvasive assay and provide a positive or negative result within 30 minutes, and the cost is less than half that of cytology. A previous study showed that the sensitivity of the NMP22 ELISA for primary BCa ranges from 44% to 100%, with a specificity of 60% to 95% [26]. Another study reported that the sensitivity in a cancer cohort (comprising patients with primary and recurrent BCa) was 40% (the sensitivity in the primary and recurrent groups alone was 42% and 34%, respectively). These results are not as good as those for the POC assay, which showed a sensitivity of 59%, 63%, and 48% in these same groups. Both assays showed a specificity of 100% in healthy individuals, while the NMP22 ELISA was 99% specific and the POC test was 93% specific in patients with benign disease [27]. A meta-analysis of 19 studies demonstrated that the pooled sensitivity and specificity of the NMP22 POC test was 56% (95% confidence interval [CI], 52%-59%) and 88% (95% CI, 87%-89%), respectively; indeed, the test showed a good ability to detect BCa in both Asian and White populations [28]. Another study compared the NMP22 biomarker with VUC and found that NMP22 was more sensitive for detecting BCa than VUC, especially among patients with microscopic hematuria (60% vs. 35%). However, the specificity of NMP22 was lower than that of VUC (78% vs. 97%, respectively) owing to the presence of NMP22 in normal urothelial cells [29]. Consequently, false-positive results are common in patients with stones, inflammation, and hematuria [30].

The BTA STAT and BTA TRAK tests target human complement factor-H related protein (hCFHrp), which is found in BCa cells and inhibits the complement cascade to prevent cell lysis. BTA STAT is a qualitative POC immunochromatographic assay, whereas BTA TRAK is a quantitative ELISA. Both have been approved by the FDA for monitoring BCa recurrence, but only as adjuncts to cystoscopy. These tests are more sensitive than VUC; a meta-analysis showed that the sensitivities of the POC and ELISA tests are 64% (95% CI, 58%–69%) and 65% (95% CI, 54%–75%), respectively [31], with specificities of 77% (95% CI, 73%–81%) and 74% (95% CI, 64%–82%), respectively. Generally, the sensitivity of the POC test ranges from 57% to 82%, with a specificity of 68% to 93% [32-34], whereas the ELISA has a sensitivity of 66% to 77% and a specificity of 50% to 75% [35,36]. However, similar to NMP22, the BTA assay also exhibits a higher false-positive rate in patients with hematuria, urolithiasis, inflammation, and other genitourinary malignancies, and in those undergoing intravesical bacille Calmette–Guérin (BCG) therapy [37].

The UroVysion bladder cancer kit is a multicolor fluorescent in situ hybridization (FISH) assay designed to estimate aneuploidy of chromosomes 3, 7, and 17, or loss of the 9p21 locus. Its performance with respect to diagnosis and surveillance for BCa has been approved by the FDA. The sensitivity of this test ranges from 69% to 87%, with a specificity between 89% and 96% [38,39]. Similarly, the sensitivity and specificity from a recent meta-analysis (11 studies) was 63% (95% CI, 50%-75%) and 87% (95% CI, 79%-93%) [31]. Another study showed that this kit detected almost twice as many NMIBC tumors as VUC, and identified 88% of invasive tumors (32% of these tumors were missed by VUC) [40]. This test is superior to the NMP22 and BTA tests owing to its high specificity. The assay is not affected by hematuria, inflammation, or other conditions that may result in false-positive readings. Thus, it could be used as an adjunct to VUC, thereby increasing sensitivity while maintaining specificity [41]. Moreover, a preponderance of evidence suggests a role for the UroVysion test for predicting responses to intravesical immunotherapy and BCG treatment [42,43].

The ImmunoCvt assay (also marketed as uCvt+) uses three fluorescently labeled monoclonal antibodies to detect high-molecular-weight forms of carcinoembryonic antigen (CEA) and two bladder tumor cell-associated mucins (LDq10 and M344) that are expressed on urothelial cells shed by tumors. This is the only commercially available test that can be used for BCa follow-up. This fluorescent test has an overall sensitivity of 40% to 100% and a specificity of 62% to 84% [44-49]. A previous case series reported a sensitivity of 74% to 87% and a specificity of 62% to 78% [50], whereas a meta-analysis of 14 studies reported a sensitivity of 78% (95% CI, 68%-85%) and a specificity of 78% (95% CI, 72%-82%) [31]. Another meta-analysis based on data from seven studies reported a pooled sensitivity and specificity of 72.5% (95% CI, 68.3%-76.5%) and 65.7% (95% CI. 62.9%-68.5%), respectively. while the pooled sensitivity and specificity of cytology were 56.6% (95% CI, 52.1%-61.1%) and 90.6% (95% CI, 88.7%-92.3%), respectively [51]. A key advantage of the ImmunoCyt fluorescent test over the NMP22, BTA, and UroVysion assays is the great improvement in sensitivity for low-grade tumors; indeed, the sensitivity increases from 63% for pTa tumors to 80% for pT1 tumors [37]. The sensitivity of the NMP22, BTA, and UroVysion assays for low-grade tumors is poor [30,52,53].

ICUROLOGY

In common with other protein-based assays, the Immuno-Cyt fluorescent test is also significantly affected by urinary tract infections, urolithiasis, and benign prostate hyperplasia, which may lead to false-positive results. The detailed characteristics of these assays are described in Table 1.

3. Non-FDA-approved urinary biomarkers for bladder cancer

Here, we summarize proposed DNA-based, RNA-based, and proteomic/peptidomic markers of BCa.

1) DNA-based urinary biomarkers

DNA tests used for surveillance usually detect loss of heterozygosity, gene methylations, and mutations in tumor cells. DNA methylation and mutations are important for the etiology and pathogenesis of many cancers [54-56], including BCa, which is a highly heterogenetic disease [57]. Thus, recent studies have explored molecular classification of both NMIBC and MIBC based on these alterations [58,59]. Alterations in DNA methylation patterns are hallmarks of cancer. Hypomethylation events may result in abnormal activation of genes, which are commonly repressed by DNA methylation. However, hypermethylation of CpG dinucleotides in the promoter regions of tumor suppressor genes can inhibit their transcription in human cells, which gives cancer cells tremendous benefits [59,60]. Thus, methylation status is one of the most studied biomarkers in the follow-up scenario; it is also used to predict treatment responses because it is both chemically stable and quantifiable in liquid biopsies [61].

The Bladder EpiCheck urine test is an in vitro diagnostic device produced by Nucleix, Ltd (San Diego, CA, USA). The test analyzes a panel of 15 DNA methylation patterns to detect BCa and has a CE mark, meaning that it is available commercially in Europe. It is effective for monitoring BCa recurrence, thereby minimizing the need for invasive cystoscopy [62]. A validation study of 222 NMIBC patients undergoing surveillance showed 90% sensitivity, 83% specificity, and a negative predictive value (NPV) of 97% [63]. In another study, Witjes et al. [62] designed a blinded, singlearm, prospective multicenter study to evaluate the performance of the EpiCheck urine test for detecting NMIBC recurrence. The overall sensitivity, specificity, and NPV were 68.2% (95% CI, 52.4%-81.4%), 88.0% (95% CI, 83.9%-91.4%), and 95.1% (95% CI, 91.9%-97.3%), respectively. Remarkably, the test could discriminate the absence of high-grade NMIBC with an NPV of 99%; by contrast, it detected the presence of high-grade NMIBC with a sensitivity of 92% [62]. D'Andrea et al. [64] published another multicenter and independent study based on data from 357 NMIBC patients. They showed that the urine test had an overall sensitivity of 67% (95% CI, 52%-80%), a specificity of 88% (95% CI, 84%-91%), and an NPV of 94.4% (95% CI, 91%-97%). The sensitivity, specificity, and NPV for high-grade and low-grade cancers were 89% (95% CI, 65%-99%) vs. 40% (95% CI, 19%-64%); 88% (95% CI, 84%–91%) vs. 88% (95% CI, 84%–91%); and 99% (95% CI, 97%-100%) vs. 96% (95% CI, 93%-98%), respectively [64]. Such consistent results make this urine test an attractive choice for use in clinical decision-making. The high NPV means that clinicians can have high confidence that a negative result rules out tumor recurrence. Accordingly, application of this test could reduce the current burden of repeat cystoscopy and cytology tests. Moreover, the results are consistent under the presence of inflammation in the urinary tract. However, the test is not simple to perform because a dedicated technician and an equipped laboratory are needed; in addition, it is expensive [62,65].

Recently, Nucleix announced the launch of its BE Safe @Home project, which provides an informatics service for NMIBC patients under surveillance with the Bladder Epi-Check urine test. In consultation with world-leading urologists, the project was implemented in Israel, Spain, and the Netherlands with a view to making surveillance more convenient during the COVID-19 pandemic and beyond.

Other markers related to methylation status are also used to follow-up BCa. Zuiverloon et al. [66] developed a methylation detection assay (based on voided urine) for specific detection of recurrence in patients with NMIBC. A logistic regression model based on methylation of a four-gene panel that combines the APC_a (APC regulator of WNT signaling pathway), TERT a (telomerase reverse transcriptase), TERT_b, and EDNRB (endothelin receptor type B) genes correlated with BCa recurrence, providing a sensitivity and specificity of 63.3% and 58.3%, respectively, in the test cohort, and of 72.3% and 55.2%, respectively, in the validation cohort [66]. Another study showed a considerably higher sensitivity and specificity (80% and 97%, respectively) using a model based on hypermethylation of SOX1 (SRY-box transcription factor 1) and IRAK3 (interleukin 1 receptor-associated kinase 3) and hypomethylation of a specific LINE1 element in MET (mesenchymal epithelial transition) in urine from BCa patients [67]. Methylation of TWIST1 (twist homolog 1) and NID2 (nidogen 2) is linked to BCa [68-70]. One study showed that under adjusted thresholds, methylation of TWIST1 and NID2 has a sensitivity and specificity of 75% and 71%, respectively [69]. More recently, a multi-institutional study reported comparable results using these two methylated genes; this study showed a sensitivity of 58% to 67% and a specificity of 61% to 69%. However, the article noted that prior BCG

BTA, bladder tumor antigen; CEA, carcinoembryonic antigen; ELISA, enzyme-linked immunosorbent assay; FDA, U.S. Food and Drug Administration; FISH, fluorescent in situ hybridization; hu-Veed trained laboratory Veed trained laboratory Veed trained laboratory High false-positive rate Veed trained laboratory High false-positive rate High false-positive rate High false-positive rate High false-positive rate -ow sensitivity in lowower specificity than -ow sensitivity in low--ow sensitivity in low--ow sensitivity in low--ower specificity than -ower specificity than -ower specificity than -ower specificity than -ow sensitivity in low--imitations grade tumor grade tumor grade tumor grade tumor grade tumor technicians technicians technicians technicians VUC VUC VUC VUC VUC Higher sensitivity than VUC (especially in low-grade Advantages High specificity Noninvasive Noninvasive Voninvasive Noninvasive Noninvasive Noninvasive tumor) Quick Refer-[38,39] [44-49] [31] [35,36] [31] [35,36] ence 31] [26] [27] [31] [51] [27] [28] [29] 50] Specificity 62.9-68.5 99-100 93-100 87–89 60-95 64–82 79-93 62-84 62-78 72-82 50-75 73-81 68-93 89-96 (%) 78 Sensitivity 44-100 40-100 58.3-76.5 74-87 68-85 34-40 52-59 54-75 58–69 57–82 69-87 50-75 48-63 66-77 (%) 60 Diagnosis, follow-up Purpose follow-up follow-up follow-up follow-up **Diagnosis**, **Diagnosis**, **Diagnosis**, Diagnosis, Follow-up Antigens/metabolites Assessed substance DNA (aneuploidies) Peptides Peptides Proteins Proteins Quantitative test: ELISA Quantitative test: ELISA Immunofluorescence Qualitative test: POC Qualitative test: POC Assay FISH Chromosomes 3, 7, and 17, Biomarker CEA, LDq10, and M344 and 9p21 hCFHrp NMP22 BTA TRAK (Polymedco) hCFHrp NMP22 BTA STAT (Polymedco) **VMP22 BladderChek** NMP22 test (Abbott) UroVysion (Abbott) mmunoCyt/uCyt+ Test name (DiagnoCure) (Abbott)

Urinary biomarker for bladder cancer

able 1. FDA-approved urinary biomarkers

man complement factor-H related protein; NMP, nuclear matrix protein; POC, point-of-care; VUC, voided urine cytology.

ICUROLOGY

treatment for NMIBC reduced the accuracy [70]. These findings are promising, despite the limited sensitivity and specificity. However, these markers were identified by research institutions with limited cohorts; thus, large validation tests and methodologic improvements are needed to achieve more accurate results.

Gene mutations are related to carcinogenesis of BCa. FGFR3 (fibroblast growth factor receptor 3) is one of the most studied genes; mutations in this gene are found in over 80% of patients with low-grade BCa and are related to a good prognosis [71]. Another well studied gene is *TERT* (telomerase reverse transcriptase), which has been investigated as a prognostic marker for NMIBC recurrence [72,73]. Allory et al. [74] found that the sensitivity of detecting NMIBC relapse was 19% for *FGFR3*, 42% for *TERT*, and 50% for *FGFR3* and *TERT* combined; for comparison, the specificities of mutations in *TERT*, *FGFR3*, and a combination of the two, were 73%, 90%, and 71%, respectively.

The Uromonitor urine-based test (Uromonitor, Porto, Portugal) is an ultra-sensitive assay capable of detecting trace amounts of TERT promoter and FGFR3 mutations in tumor cells shed into urine [75]. Screening of targeted alterations is based on a highly sensitive multiplex competitive allelespecific discrimination PCR that allows clear interpretation of results. Compared with Sanger sequencing, this test can detect a very small number of altered cells in a large pool of unaltered cells. In addition, it is superior to next generation sequencing (NGS)-based assays in terms of cost and time. A multicenter validation study revealed that the Uromonitor urine-based test has a sensitivity of 73.5% and a specificity of 73.2% for detection of NMIBC recurrence [76]. Higher sensitivity and specificity were reported in a study based on a cohort of 72 patients first diagnosed with BCa and under surveillance for NMIBC; the data suggest that the sensitivity, specificity, and NPV of the Uromonitor urine-based test are 100%, 96.3%, and 88.9%, respectively [75]. Concerning lowgrade and high-grade recurrence-positive patients, the test showed a detection rate of 625% and 75%, respectively. Also, the presence of inflammation or other benign lesions in the urinary tract did not affect results. Thus, routine use of this urine-based test plus cystoscopy could be very cost-effective. Indeed, the sensitivity and specificity are similar to those of cystoscopy (which were 79.4% and 73.2%, respectively) in one study, which suggests that the Uromonitor urine-based test is an appropriate option when cystoscopy cannot be performed or is not available routinely [76].

The Uromonitor-V2 urine-based assay (Uromonitor) added the *KRAS* (kirsten rat sarcoma 2 viral oncogene homolog) hotspot mutation to the Uromonitor kit. The Uromonitor-V2 assay showed 100% sensitivity, 833% specificity, and an NPV of 100% for evaluating NMIBC recurrence in a multicenter study of 122 patients [76]. Another recent study reported a sensitivity, specificity, and NPV of 93.1%, 85.4%, and 95.3%, respectively, for detecting BCa recurrence, whereas VUC showed a sensitivity, specificity, and NPV of 26.3%, 90.9%, and 68.2%, respectively. Thus, the Uromonitor-V2 urine-based assay is a promising test for surveillance of BCa [77].

The UroSEEK urine-based molecular assay is another noninvasive commercially available test (although not approved by the FDA or the European organization). The assay is designed to detect alterations in 11 genes, including 10 typical mutations associated with BCa (FGFR3, TERTp, TP53 [tumor protein p53], ERBB2 [erb-b2 receptor tyrosine kinase 2], CDKN2A [cyclin-dependent kinase inhibitor 2A], KRAS, HRAS [v-ha-ras harvey rat sarcoma viral oncogene homolog], MET, PIK3CA [phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha], MLL [mixed-lineage leukemia 1], and VHL [von Hippel-Lindau tumor suppressor]) plus detection of an uploidy [78]. Springer et al. [78] found that this test could detect recurrence with a sensitivity of 68% and a specificity of 80%. Another study revealed that in the setting of early detection, the sensitivity and specificity were 96% and 88%, respectively, with an NPV of 99%; however, the results from a surveillance cohort were less robust (sensitivity of 74%, specificity of 72%, and NPV of 53%) [79]. Nevertheless, the UroSEEK molecular assay was more sensitive than cytology both in the surveillance cohort (71% vs. 25%, respectively) and in the primary detection cohort (95% vs. 43%, respectively); however, the specificity of cytology was superior in the detection cohort (100% vs. 93%) [78]. Accordingly, this test does not show excellent performance for the follow-up of patients with a prior diagnosis of BCa.

All these findings strongly suggest that mutations or methylation status of several genes are promising biomarkers for BCa; thus, a combination of genetic and epigenetic markers for BCa diagnosis and surveillance is both logical and appealing.

The combination of FGFR3 mutations and methylation biomarkers has been tested, with promising results. Beukers et al. [80] investigated the performance of FGFR3 and TERTmutations combined with OTX1 (orthodenticle homeobox 1) methylation in 977 patients with NMIBC. They reported that the sensitivity for detecting NMIBC recurrence was 57%, with a specificity of 59% [80]. Similarly, a 3-plex methylation (combination of OTX1, ONECUT2 [one cut homeobox 2], and OSR1 [odd-skipped related transcription factor 1] methylation) assay combined with the FGFR3 mutation assay detects recurrent NMIBC in voided urine with a sen-

sitivity of 79% and a specificity of 77% [81]. Another study combined *FGFR3* mutation with methylation of a set of DNA markers (*HS3ST2* [heparan sulfate-glucosamine 3-sulfotransferase 2], *SEPTIN9*, and *SLIT2* [slit guidance ligand 2]) and reported a sensitivity of 94.5% for discriminating recurrent tumors. The specificity and NPV for this assay were 75.9% and 98.5%, respectively [82]. Recently, MDxHealth SA (Euronext: MDXH.BR; Herstal, Belgium) announced the commercial launch of its AssureMDx laboratory-based test in the United States, which combines methylation (*OTX1*, *ONECUT2*, and *TWIST1*) and mutation (*FGFR3*, *TERT*, and *HRAS*) biomarkers to identify BCa in patients with hematuria [83]. A multicenter study verified a sensitivity of 93% and a specificity of 86% for BCa diagnosis [84].

Thus, combined analyses of DNA mutations and DNA methylation markers could be used for risk stratification of patients with BCa and for surveillance, forming the foundation for a promising noninvasive urine test.

2) RNA-based urinary biomarkers

RNA-based urinary biomarkers are less well studied than DNA markers. One commercially available RNA test is the Cxbladder Monitor (Pacific Edge Diagnostics, Dunedin, New Zealand). This test evaluates the expression of five urinary mRNAs (insulin-like growth factor binding protein 5 [IGFBP5], homeobox protein Hox-A13 [HOXA13], midkine [MDK], cyclin-dependent kinase 1 [CDK1], and chemokine receptor type 2 [CXCR2]) and incorporates them into a mathematical algorithm that also includes clinical variables, such as primary vs. recurrent BCa and time since tumor occurrence, to generate a score to give a positive or negative result [85]. Lotan et al. [85] compared the Cxbladder Monitor with current FDA-approved urine tests; they examined 1,036 urine samples from 803 patients undergoing surveillance for recurrent BCa and found that the Cxbladder Monitor test significantly outperformed the other tests. The Cxbladder Monitor showed a sensitivity of 91% (95% CI, 86%-95%) when monitoring a BCa population. This is obviously higher than that of other tests: 22% for cytology, 26% for the NMP22 ELISA test, and 11% for NMP22 BladderChek. The NPV of the Cxbladder Monitor was also superior (96%) to that of cytology (87%), the NMP22 ELISA (87%), and NMP22 BladderChek (86%) [85]. In addition, the Cxbladder Monitor showed a sensitivity of 93% (95% CI, 85%-97%) and an NPV of 94% (95% CI, 88%-97%) in patients undergoing routine surveillance for recurrent BCa [85,86]. By contrast, cytology, the NMP22 ELISA, and NMP22 BladderChek showed sensitivities of 22%, 29%, and 8%, and NPVs of 83%, 83%, and 81%, respectively. After seeing the evidence, New Zealand's public healthcare providers have integrated the Cxbladder Monitor into their routine clinical surveillance of BCa patients. A recent study demonstrated that the Cxbladder Monitor accurately detected about 77.8% of recurrence-free patients per year who could avoid unnecessary cystoscopy [87]. During the COVID-19 lockdown, two of New Zealand's public healthcare providers started using the Cxbladder Monitor for in-home testing as an out-patient solution for BCa monitoring; this is especially useful for older patients at high risk for COVID-19.

Another commercially available RNA test is the Xpert Bladder Cancer Monitor, which measures five mRNAs (v-abl Abelson murine leukemia viral oncogene homolog 1 [ABL1], corticotropin-releasing hormone [CRH], insulin-like growth factor 2 [IGF2], uroplakin 1B [UPK1B], and annexin A10 [ANXA10]) that are frequently overexpressed in BCa. This test provides qualitative monitoring of BCa recurrence within 90 minutes. Wallace et al. [88] developed this urine-based test using 450 urine specimens collected from 18 multinational sites and obtained an overall sensitivity of 73%, with specificities of 90% and 77% in hematuria and surveillance patient populations, respectively. In another study, Pichler et al. [89] examined 140 patients with a history of NMIBC who were undergoing routine surveillance and reported for the first time that the Xpert Bladder Cancer Monitor outperforms VUC in terms of sensitivity (84% vs. 33%, respectively) and NPV (93% vs. 76%, respectively), even in those with lowgrade and Ta tumors; however, the specificity of the two tests was similar (91% vs. 94%, respectively). Another multicenter study compared the Xpert Bladder Cancer Monitor, VUC, and the UroVysion bladder cancer kit to determine their follow-up performance in patients previously diagnosed with NMIBC. The Xpert Bladder Cancer Monitor showed a higher sensitivity and NPV (74% [95% CI, 94%-99%] and 93% [95% CI, 89%–96%]), respectively, than did VUC or the UroVysion bladder cancer kit. Moreover, the sensitivity and NPV for high-grade tumors were 83% (95% CI, 64%–93%) and 98% (95% CI, 94%-99%), respectively [90]. The improved NPV of this test in patients under follow-up for BCa suggests that the Xpert Bladder Cancer Monitor is a promising tool for excluding BCa and reducing the need for cystoscopy. However, D'Elia et al. [91] indicated a lower overall sensitivity: 46.2% for detecting NMIBC recurrence. In addition, results from the Bladder Cancer Italian Active Surveillance project, which enrolled 106 patients with low-grade NMIBC who developed recurrence during follow-up and underwent active surveillance, suggest the need to optimize the cutoff value [92]. Thus, further research on larger populations is mandatory before this test can be used routinely in clinical

practice.

In addition to these mRNA-based biomarkers, a number of urinary microRNA (miRNA) biomarkers are emerging. miRNAs interact with their target mRNAs to modulate their expression, thereby controlling many physiologic processes, including carcinogenesis [93]. Most miRNA-based studies have focused on the diagnostic performance of the miRNAs (one special miRNA or miRNA panels) that are differentially expressed in BCa urine [8,94,95]. As a prognostic marker for predicting NMIBC recurrence. Kim et al. [96] found that urinary miR-214 was down-regulated in NMIBC patients who experienced recurrence during surveillance. with a hazard ratio of 2011 (95% CI, 1027-3.937), when compared with those without recurrence. Sapre et al. [97] examined the potential of a urinary miRNA panel for predicting the presence of BCa in NMIBC patients undergoing surveillance. They found that a combination of six miRNAs (miR16, miR200c, miR205, miR21, miR221, and miR34a) vielded an area under the curve (AUC) of 0.85 for distinguishing NMIBC patients with recurrence from those without in the discovery cohort, and they showed high sensitivity (88%) and adequate specificity (48%) (AUC=0.74) in the validation cohort; these data suggest that cystoscopy rates in the validation cohort would have been reduced by 30%.

Lately, another group of noncoding RNAs, named long noncoding RNAs (IncRNAs), which are longer than miRNAs, have emerged as an informative tool for the management of BCa [98]. The advent of high-throughput technology, such as RNA-seq, has identified more than 10,000 unique lncRNAs and clarified their biological functions. lncRNAs play a crucial role in BCa tumorigenesis by modulating cellular pathways involved in cell transformation [99-101]. Zhang et al. [102] investigated the potential application of an lncRNA called urothelial cancer associated 1 (UCA1), which is found in BCa patients' urine. They identified that UCA1 showed high sensitivity and specificity (84.4% and 92.4%, respectively) for BCa (AUC=0.898). Moreover, the study highlighted the role of UCA1 as a prognostic biomarker for NMIBC patients who may progress to MIBC (sensitivity, 86.4%; specificity, 92.3%) [102]. Similarly, Eissa et al. [103] showed that UCA1 has great sensitivity (91.5%) and specificity (96.5%) for detecting BCa. Indeed, urinary UCA1 was more accurate than VUC in NMIBC patients [103]. Therefore, accumulation of UCA1 in urine may be a prospective marker for BCa diagnosis and surveillance. HOX antisense intergenic RNA (HOTAIR) is a recently discovered lncRNA that plays an important role in BCa. A study revealed that HOTAIR expression has prognostic value for BCa progression, recurrence, and survival [104]; recurrent NMIBC tumors showed significantly higher

ICUROLOGY

HOTAIR expression than did nonrecurrent tumors. Kaplan-Meier analysis revealed that patients with higher HOTAIR expression exhibited significantly earlier recurrence and earlier progression after recurrence. Another pilot study detected lncRNA H19 in urine sediment from 90.5% of BCa patients and 25.9% of healthy controls, making it a supplemental tool for BCa diagnosis [105]. Although expression of other lncRNAs correlates with BCa, most studies were based on tissues or cell lines; therefore, the results require validation in urine samples to confirm their practical applicability as noninvasive BCa biomarkers [106-108]. In addition, urinebased research mostly relies on urine sediment, and different lncRNA targets are detected in urine cells. Thus, results may vary because of the paucity of genitourinary-derived cells in urine [109,110], the presence of urinary crystals [111], and the concentration of inhibitors in urine sediment [112,113]. A recent study reported a 16 cell-free urinary lncRNAbased panel, which showed differential expression between NMIBC and urocystitis patients [114]. Among them, the AUCs for four biomarkers (UCA1-201, HOTAIR, HYMA1, and metastasis associated lung adenocarcinoma transcript 1 [MALAT1]) were higher than 0.80, suggesting superior diagnostic performance in differentiating NMIBC from urocystitis. Next, machine learning was used to train the four lncRNA panel as a predictive model; the panel made good predictions in the validation phase, showing a sensitivity of 93.3% and a specificity of 96.7% for discriminating NMIBC from urocystitis.

3) Proteomic/peptidomic urinary biomarkers

The UBC urinary bladder cancer test (IDL Biotech, Borläbger, Sweden) is a commercially available test that measures soluble fragments of cytokeratin 8 and 18 (CK8 and CK18) in urine samples for the purpose of diagnosis and monitoring of BCa. The UBC test is available in ELISA or POC formats. The sensitivity and specificity vary from 12% to 88% and from 77% to 92%, respectively [115]. Pichler et al. [116] found a sensitivity of 61.3% and 64.5%, and a specificity of 77.3% and 81.8%, for the qualitative and quantitative UBC tests, respectively; thus, the assay is more sensitive than VUC (sensitivity, 25.8%) and NMP22 (sensitivity, 12.9%). A multicenter study demonstrated another interesting outcome: the sensitivity for high-grade NMIBC was greater than that for low-grade NMIBC (75.0% vs. 38.8%, respectively), suggesting the potential of UBC as a clinically valuable urinary protein biomarker for detection of high-grade NMIBC [117]. Babjuk et al. [118] reported that the sensitivity and specificity of the UBC ELISA test were 12.1% and 97.2%, respectively, for detecting NMIBC in patients under

surveillance. However, when setting a new cutoff to reach a sensitivity of 90%, the specificity declined to 20.4%, indicating that individually installed cutoffs are of no benefit [118]. Concerning the issue of an appropriate cutoff for the UBC test, a study compared the uncorrected and corrected cutoff values of the UBC test for distinguishing BCa patients with and without a recurrence. The overall sensitivity, specificity, and NPV of the uncorrected UBC test were 20.7%, 84.7%, and 72.6%, respectively; those for the corrected UBC test were 20.7%, 79.2%, and 71.3%, respectively. Nonetheless, the receiver operating characteristic analyses showed no statistical significance (both p>0.05), indicating that the UBC test has no diagnostic value [119]. In contrast to other markers, the UBC test is rapid, with results available within 10 minutes; however, the clinical utility of the UBC test for followup of BCa patients remains unconvincing.

The ADXBLADDER in vitro diagnostic test (Arquer Diagnostics, Sunderland, UK) is an up-to-date commercial urine test that detects mini-chromosome maintenance complex component 5 (MCM5), which is a marker of cells that are replicating (or that still have the capability to replicate). In July 2020, the UK National Health Service approved the use of the ADXBLADDER test to help with diagnosis and surveillance of BCa [11]. The test was superior to VUC for detecting BCa recurrence [120]. The ADXBLADDER MCM5 test has sensitivities of 44.1% and 58.8% for low- and highgrade recurrence, respectively, which is more accurate than VUC (sensitivity of 17.6% for both low-grade and high-grade recurrence); thus, this test could be a reliable alternative to VUC for follow-up monitoring. Similar results were reported in a multicenter prospective, blinded study carried out from August 2017 to July 2019 at 21 European Union centers. The study demonstrated that the ADXBLADDER test excluded the presence of high-grade recurrence in 97.8% of cases (compared with 97.1% with VUC). Meanwhile, the sensitivity of the test was 51.9%, which was much higher than that of VUC (16.7%) [121]. The test demonstrated an impressive NPV of 92% to 99%, using a standard ELISA and with a rapid 2-hour turnaround time. Despite having advantages over VUC, the overall performance of the ADXBLADDER test remains relatively low. The clinical implementation of these biomarkers for the follow-up of BCa must be investigated further in prospective randomized trials in patients with low-grade as well as high-grade tumors.

Cytokeratin fragment 21.1 (CYFRA 21.1) is an ELISAbased assay that measures the concentration of a soluble fragment of cytokeratin 19, the levels of which in urine samples differ between healthy persons and those with BCa (sensitivity, 82%; specificity, 80%) [122]. An extensive metaanalysis study reported not only that the CYFRA 21.1 level has diagnostic value for BCa, but also that CYFRA 21.1 levels are higher in those with metastatic BCa than in those with locally invasive disease, inferring a role for detecting metastases [123] Moreover, Nisman et al. [124] showed that CYFRA 21.1 detected 100% of carcinoma *in situ* cases, 92.8% of MIBC cases, and 91.9% of grade 3 tumors; also, the assay detected 65% of recurrent tumors and 71% of primary tumors missed by VUC. Unfortunately, like other protein markers in urine, CYFRA 21.1 has a high false-positive rate in patients with urinary tract infections, stones, or a history of pelvic radiotherapy, urethral catheterization, or BCG intravesical instillation within the 3 previous months.

The characteristics of non-FDA-approved but commercially available biomarkers are shown in Table 2.

4. Limitations of current urinary biomarkers

Although newly developed molecular biomarkers for BCa management show superior sensitivity to VUC, few are included in clinical guidelines. Most studies concur that the currently FDA-approved biomarkers do not perform well in real clinical practice as they lack sensitivity or specificity for detecting BCa, or the studies suffered from high false-positive rates. Indeed, a systemic meta-analysis of 57 studies concluded that the false-positive rates of FDA-approved markers range from 44% to 78% [31]. This review also showed that these urinary biomarkers missed 18% to 43% of patients with BCa. These poor performances suggest that limited sensitivity, specificity, and high false-positive rates are the greatest challenge to application of urinary biomarkers in clinical practice.

Most studies of these markers reported "promising" results and initially "positive" observations, which may contribute to early clinical application. However, these preliminary findings lose their glory when tested in real clinical practice; such differences in performance may be related to study design (i.e., selection of suitable cohorts, sufficient number to achieve statistical significance, inadequate followup, and poor validation of results). Moreover, the definition of an ideal marker contains the words easier, faster, better, and cheaper [125]. Easier allows performance in a clinical environment; faster means a rapid turnaround; better means that it must at least be equal to currently clinically available alternatives and provide information that is helpful to clinicians with respect to management of the disease; and cheaper means a reduced economic burden on patients and health services. Nevertheless, many urine tests require highly skilled personnel and specialized laboratory equipment, which increases both time and cost. Initially, the cost of

lable 2. Commercially available (non-FUA-approved) urine tests for bladder cancer	e (non-FUA-approved	I) urine tests	tor bladder ca	ncer						
Test name	Biomarker	Assay	Assessed substances	Purpose	Sensitivity (%)	Specificity (%)	NPV	Refer- ences	Advantages	Limitations
DNA-based assays Bladder EpiCheck (Nucleix)	15 DNA methylation	Real-time PCR	Urinary tumor Follow-up DNA	Follow-up	Overall: 67–90 High-grade: 89–92 Low-grade: 40	Overall: 83–88 High-grade: 88 Low-grade: 88	Overall: 94.4–97 High-grade: 99 Low-grade: 96	[62-64]	Noninvasive Higher sensitivity than VUC High NPV Provide in-home testing service	Low sensitivity in low-grade tumor Lower specificity than VUC High false-positive rates Need a dedicated technician and an equipped laboratory
Uromonitor (Uromonitor)	DNA [*] mutations	Real-time PCR	Urinary tumor DNA	Follow-up	Overall: 73.5–100 High-grade: 75 Low-grade: 62.5	Overall: 73.2–96.3	Overall: 88.9	[76,77]	Noninvasive Higher sensitivity than VUC High NPV Lower false-positives than other tests	Low sensitivity in low-grade tumor Lower specificity than VUC Need a dedicated technician and an equipped laboratory
Uromonitor-V2 (Uromonitor)	DNA ^a mutations	Real-time PCR	Urinary tumor DNA	Follow-up	Overall: 93.1–100	Overall: 83.3–85.4	Overall: 95.3-100	[77,78]	Noninvasive Higher sensitivity than VUC High NPV No confusions from other benign diseases	Need a dedicated technician and an equipped laboratory
UroSEEK	10 gene mutations ^b	NGS	Urinary tumor DNA	Diagnosis, follow-up	95–96 68–74	88–100 72–80	23 66	[08'62]	Noninvasive Higher sensitivity than VUC (especially for early detec- tion of BCa) High specificity and NPV for BCa early detection	Lower sensitivity, specificity and NPV for BCa follow-up surveil- lance Need a dedicated technician and an equipped laboratory Complicated methods
AssureMDx (MDxHealth) RNA-based assays	DNA ^c methylation+ DNA ^d mutation	Real-time PCR	Urinary tumor DNA	Diagnosis	93	86	NA	[85]	Noninvasive Higher sensitivity than VUC	Need more validation results Need a dedicated technician and an equipped laboratory
Cx Bladder Monitor (Pacific Edge Diagnostics)	5 mRNA [®] expression	Real-time PCR	Urinary tumor RNA	Diagnosis, follow-up	91 93	85	96 94	[86,87]	Noninvasive Higher sensitivity Higher NPV Provide in-home testing service	Need more validation results Need a dedicated technician and an equipped laboratory
Xpert Bladder Cancer Monitor	5 mRNA ^ŕ expression	Real-time PCR	Urinary tumor Follow-up RNA	Follow-up	Overall: 46.2–84 High-grade: 83	90-91	Overall: 77–93 High-grade: 98	[89-92]	Noninvasive Qualitative monitoring of BCa recurrence in 90 minutes Higher sensitivity than VUC (Especially in high-grade tumor)	Need an optimization of its cutoff Need further research on larger population

Table 2. Commercially available (non-FDA-approved) urine tests for bladder cancer

ICUROLOGY

Table 2. Continued

inutes r detecting a other	Test name	Biomarker	Assay	Assessed substances	Purpose	Sensitivity (%)	Sensitivity Specificity (%) (%)	NPV	Keter- ences	Advantages	Limitations
MCM5 ELISA Peptide Diagnosis, High-grade: NA 92–99 [121,122] Noninvasive follow-up 51.9–58.8 Superior to VUC for detecting Low-grade: A.1 No confusions from other 44.1 benian diseases	Proteomic/Peptidomic assays UBC Rapid and ELISA (IDL Biotech)	CK8, CK18	POC/ELISA	Peptide	Diagnosis, follow-up	12–88 12.1	77–92 97.2	AN	[117,119]	Noninvasive Quick test in 10 minutes	Need an optimization of its cutoff Need further research on larger population Clinical utility in the follow-up needs more convincing results
	ADXBLADDER (Arquer Diagnostics)	MCM5	ELISA	Peptide	Diagnosis, follow-up	High-grade: 51.9–58.8 Low-grade: 44.1	NA	92-99	[121,122]	Noninvasive Superior to VUC for detecting recurrence of BCa No confusions from other benign diseases	Inadequate sensitivity for clinical utility Need further research on larger population

corticotropin-releasing hormone; CXCR2, chemokine receptor type 2; ELISA, enzyme-linked immunosorbent assay; ERBB2, erb-B2 receptor tyrosine kinase 2; FDA, U.S. Food and Drug Administration; *FGFR3*, fibroblast growth factor receptor 3; HOX413, homeobox protein Hox-A13; HRA5, v-ha-ras harvey rat sarcoma viral oncogene homolog; *IGF2*, insulin-like growth factor 2; *IGFBP5*, insulin-like transition; MLL, mixed-lineage leukemia 1; NA, not available; NGS, next generation sequencing; NPV, negative predictive value; ONECUT2, one cut homeobox 2; OTX1, orthodenticle homeobox 1; 4BL1, v-abl Abelson murine leukemia viral oncogene homolog 1; ANXA10, annexin A10; BCa, bladder cancer; CDK1, cyclin-dependent kinase 1; CDKN2A, cyclin-dependent kinase inhibitor 2A; CRH, growth factor binding protein 5; KR45, kirsten rat sarcoma 2 viral oncogene homolog; MCM5, mini-chromosome maintenance complex component 5; MDK, midkine; MET, mesenchymal epithelial PCR, polymerase chain reaction; *PIK3CA*, phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha; POC, point-of-care; *TERT*, telomerase reverse transcriptase; *TP53*, tumor protein p53; *WIST1*, twist homolog 1; VHL, von Hippel-Lindau syndrome; VUC, voided urine cytology; *UPK1B*, uroplakin 1B.

:TERT promoter and the FGFR3.

"FGFR3, TP53, CDKN2A, ERBB2, HRAS, KRAS, PIK3CA, MET, VHL, MLL, and TERT promoters.

:OTX1, ONECUT2, and TWIST1

:FGFR3, TERT, and HRAS.

:IGFBP5, HOXA13, MDK, CDK1, and CXCR2. ABL 1, CRH, IGF2, UPK1B, and ANXA 10.

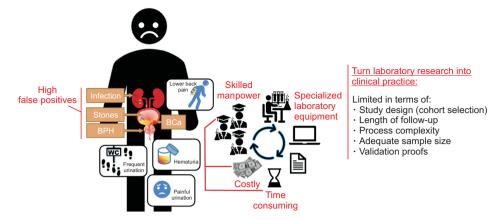


Fig. 2. Limitations of current urine tests. Lack of disease-specific symptoms means that BCa can be confused with other benign diseases such as infection, stones, and BPH, all of which can cause false-positive results in tests based on BCa diagnostic markers. Currently developed markers are more suitable for laboratory research than for use in clinical practice. There are several reasons for this. Studies in the field of biomarker research are limited with respect to appropriate cohort selection. Also, recent studies comprise too few samples, and follow-up is too short for meaningful statistical analysis. Additionally, laboratory-derived biomarkers are still relatively new and need to be validated in larger cohorts. Moreover, tests and assays often require skilled technicians and specialized equipment; thus, they are time-consuming and expensive. Although the biomarker market has grown rapidly, many problems await solutions. BCa, bladder cancer; BPH, benign prostatic hyperplasia; WC, water closet.

laboratory research is covered by the sponsor of the research project; however, these costs are then passed on to patients and hospitals when the tests enter clinical use.

Owing to the lack of disease-specific symptoms, diagnosis and follow-up of BCa are challenging. Noninvasive urine tests are designed to diagnose BCa, but it remains unclear how they can be integrated effectively into clinical decisionmaking, particularly with respect to discrimination of BCa from other diseases presenting with similar signs and symptoms. This pertains largely to patients with gross hematuria, non-urinary tract infection-related irritative voiding symptoms, and those with microscopic hematuria found on routine urinalysis. These variables should be taken into account during study design, although they are often neglected. The majority of studies are case-control trials comparing artificially composed study cohorts, in which the prevalence of the cancer frequently exceeds 50%. This high disease prevalence is not common in urological practice, and such evaluations lead to an overly optimistic calculation of positive predictive value. Finally, poor study design with respect to selection of patient cohorts and endpoints, and statistical considerations, is one of the reasons for the limited incorporation of novel BCa urinary markers into clinical decision-making. This may be connected to the lack of coherent and comprehensive processes (pipelines) for biomarker development. Therefore, to improve and standardize BCa marker development, Goebell et al. [126] proposed a stepwise procedure comprising four phases (analogous to therapeutic trials). Shariat et al. [127] placed emphasis on this four-phase process and stressed the importance of statistical considerations when conducting research into clinical biomarkers for BCa. It is crucial that researchers are aware of the complexity and poor success rate of biomarkers trying to enter the clinical arena. Fig. 2 describes the limitations of the current urine tests.

THE MICROBIOME AS A SOURCE OF BIOMARKERS

Mounting evidence indicates that the microbiota plays an important role in carcinogenesis and response to treatment. The dogma that urine is sterile has been discredited. and dysbiosis of the urinary microbiota is linked to urological disorders [128-132]. Developed techniques, including highthroughput sequencing, single-cell transcriptomics, and mass spectrometry, offer precise characterization of single enteric, neoplastic, and immune cells, and culture assays enable detection of microbes throughout the urinary system [131,133-135]. A recent study demonstrated an innovative mechanism underlying intravesical immunotherapy with BCG, a live attenuated strain of Mycobacterium bovis. The study suggests that the effects of BCG on BCa may be related to the action of certain microbiomes. The local microbiota may inactivate BCG directly in the bladder or may modulate urothelial sensitivity to BCG through competitive binding to fibronectin [136]. All these advances have focused BCa research on an attempt to understand the relationship between the commensal urinary microbiome and BCa development, as well as its impact on treatment efficacy through modulation of the anti-cancer immune response. The exact nature and role of the most applicable microbes remain unclear, but

their potential involvement in BCa is apparent. A study by Xu et al. [137] undertook 16S sequencing analysis of voided urine from six healthy adults and eight BCa patients. They reported that the genus Streptococcus was enriched in the BCa patients [137]. Similarly, another study compared voided urine from 12 BCa patients with that from 11 healthy adults using 16S sequencing. They identified a known colorectal cancer-related genus, Fusobacterium, and showed that it was enriched in the BCa group; however, they found no significant differences in microbial diversity or overall microbiome composition between the groups [138]. Wu et al. [139] compared 31 male BCa patients with 18 healthy control subjects using 16S sequencing of midstream voided urine; they found that enrichment of the genera Acinetobacter, Anaerococcus, and Sphingobacterium was associated with BCa. In addition, greater bacterial richness was also present in urine from NMIBC patients with a high risk for recurrence or progression, indicating that higher bacterial richness may be a potential indicator of NMIBC prognosis. More recently, Chipollini et al. [140] performed 16S ribosomal RNA sequencing in voided urine samples from 10 noncancerous controls, 12 NMIBC patients, and 15 MIBC patients and discovered that the noncancerous group had less species variation and phylogenetic diversity than both the NMIBC and MIBC groups (all p<0.05). The authors also found that Bacteroides, Lachnoclostridium, and Burkholderiaceae species were significantly enriched in noncancerous samples, whereas Bacteroides and Faecalibacterium species were enriched in MIBC samples [140].

Unfortunately, these studies have a critical limitation in that they all analyzed the microbiome in voided urine. Voided urine is not representative of the bladder microbiome because the bacterial DNA detected in midstream voided urine differ substantially from the DNA detected in transurethral catheterized urine, which may be free from external contamination [141,142]. Moreover, although 16S sequencing is a powerful and sensitive tool for microbiome research, there are limitations. First, the results of 16S rRNA gene sequencing are relative rather than absolute; this is because the actual quantity of a particular bacteria is uncertain. Second, the outcomes can be biased due to fluctuating PCR amplification frequency and incomplete reference databases used for sequence analysis. Third, it does not determine cause-andeffect relationships. Fourth, it cannot determine whether bacteria were alive or dead, thereby requiring further urine culture for confirmation [143]. Connections between the urinary microbiome and BCa seemed real, and the possibility of using it as a noninvasive biomarker is highly intriguing. Further investigations should be conducted to improve the studies in this intriguing area.

CONCLUSIONS

Urinary biomarkers have been disregarded because of a perceived lack of sensitivity, a high rate of false positivity, and a paucity of independent validation studies, even though they have great potential as biomarkers of BCa owing to their noninvasive and easy sampling methods. For this reason, substantial improvements in this area have been made in the past few years. In addition, the current COVID-19 pandemic, which has caused inevitable delays in diagnosis, has highlighted the value of quick, efficient, easy, and contactless methods that can be used in clinical practice. Thus, much effort should be devoted to translating potential urinary biomarkers into clinical practice to alleviate the backlog of patients awaiting diagnostic procedures. However, current screening and surveillance urine tests are hampered by a low disease prevalence, which results in a high cost-tobenefit ratio. Although preliminary reports suggest that urinary biomarkers are feasible, there is insufficient detailed information from large study populations to make a final judgment. We urge the scientific and clinical communities to join hands to tackle these problems by modifying protocols and conducting prospective trials to provide a basis for integrating molecular markers into clinical practice. Guidelines on the use of biomarkers would also be welcome. Also, there should be direct head-to-head comparisons of urinary biomarkers to determine the best combination that provides the greatest sensitivity and specificity. The limitations of urinary biomarkers are solvable, albeit time-consuming; however, such markers will be of great benefit to BCa patients around the world. Accordingly, we can use them to improve care for patients with BCa during the COVID-19 pandemic and beyond.

CONFLICTS OF INTEREST

The authors have nothing to disclose.

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AUTHORS' CONTRIBUTIONS

Research conception and design: Xuan-Mei Piao. Data acquisition: Xuan-Mei Piao and Howon Kang. Statistical analysis: Xuan-Mei Piao. Drafting of the manuscript: Xuan-Mei Piao. Critical revision of the manuscript: Wun-Jae Kim and Seok Joong Yun. Obtaining funding: Howon Kang and Seok Joong Yun. Approval of the final manuscript: Wun-Jae Kim and Seok Joong Yun.

DATA AVAILABILITY STATEMENT

The data presented in this study are available from the corresponding author upon reasonable request.

REFERENCES

- Wong MCS, Fung FDH, Leung C, Cheung WWL, Goggins WB, Ng CF. The global epidemiology of bladder cancer: a joinpoint regression analysis of its incidence and mortality trends and projection. Sci Rep 2018;8:1129.
- Siegel RL, Miller KD, Jemal A. Cancer statistics, 2020. CA Cancer J Clin 2020;70:7-30.
- Saginala K, Barsouk A, Aluru JS, Rawla P, Padala SA, Barsouk A. Epidemiology of bladder cancer. Med Sci (Basel) 2020;8:15.
- Meeks JJ, Al-Ahmadie H, Faltas BM, Taylor JA 3rd, Flaig TW, DeGraff DJ, et al. Genomic heterogeneity in bladder cancer: challenges and possible solutions to improve outcomes. Nat Rev Urol 2020;17:259-70.
- 5. Dagogo-Jack I, Shaw AT. Tumour heterogeneity and resistance to cancer therapies. Nat Rev Clin Oncol 2018;15:81-94.
- Soria F, Krabbe LM, Todenhöfer T, Dobruch J, Mitra AP, Inman BA, et al. Molecular markers in bladder cancer. World J Urol 2019;37:31-40.
- Flaig TW, Spiess PE, Agarwal N, Bangs R, Boorjian SA, Buyyounouski MK, et al. Bladder cancer, version 3.2020, NCCN clinical practice guidelines in oncology. J Natl Compr Canc Netw 2020;18:329-54.
- Piao XM, Jeong P, Kim YH, Byun YJ, Xu Y, Kang HW, et al. Urinary cell-free microRNA biomarker could discriminate bladder cancer from benign hematuria. Int J Cancer 2019;144:380-8.
- Goldberg H. EAU 2020: urinary markers in low-grade nonmuscle invasive bladder cancer: ready to stop cystoscopies [Internet]. San Francisco: UroToday; 2020 Jul 17–19 [cited 2021 Feb 26]. Available from: https://www.urotoday.com/ conference-highlights/eau-2020/bladder-cancer/123171-eau-2020-urinary-markers-in-low-grade-non-muscle-invasive-

 $bladder\-cancer\-ready\-to\-stop\-cystos\-copies.html.$

- Quaedackers JSLT, Stein R, Bhatt N, Dogan HS, Hoen L, Nijman RJM, et al. Clinical and surgical consequences of the COVID-19 pandemic for patients with pediatric urological problems: statement of the EAU guidelines panel for paediatric urology, March 30 2020. J Pediatr Urol 2020;16:284-7.
- Ng K, Vinnakota K, Sharma A, Kelly J, Dasgupta P, Vasdev N. Urinary biomarkers to mitigate diagnostic delay in bladder cancer during the COVID-19 era. Nat Rev Urol 2021;18:185-7.
- 12. Vaidyanathan K, Vasudevan DM. Organ specific tumor markers: what's new? Indian J Clin Biochem 2012;27:110-20.
- Babjuk M, Burger M, Compérat EM, Gontero P, Mostafid AH, Palou J, et al. European Association of Urology guidelines on non-muscle-invasive bladder cancer (TaT1 and carcinoma in situ) - 2019 update. Eur Urol 2019;76:639-57.
- Batista R, Vinagre N, Meireles S, Vinagre J, Prazeres H, Leão R, et al. Biomarkers for bladder cancer diagnosis and surveillance: a comprehensive review. Diagnostics (Basel) 2020;10:39.
- Ru Y, Dancik GM, Theodorescu D. Biomarkers for prognosis and treatment selection in advanced bladder cancer patients. Curr Opin Urol 2011;21:420-7.
- Temilola DO, Wium M, Coulidiati TH, Adeola HA, Carbone GM, Catapano CV, et al. The prospect and challenges to the flow of liquid biopsy in Africa. Cells 2019;8:862.
- Vaidyanathan R, Soon RH, Zhang P, Jiang K, Lim CT. Cancer diagnosis: from tumor to liquid biopsy and beyond. Lab Chip 2019;19:11-34.
- Di Meo A, Bartlett J, Cheng Y, Pasic MD, Yousef GM. Liquid biopsy: a step forward towards precision medicine in urologic malignancies. Mol Cancer 2017;16:80.
- Poulet G, Massias J, Taly V. Liquid biopsy: general concepts. Acta Cytol 2019;63:449-55.
- Chan JYH, Wang Z. Tumor markers. In: Lau WY. Hepatocelluar carcinoma. Singapore: World Scientific Publishing Co; 2008;159-82.
- Urquidi V, Goodison S, Ross S, Chang M, Dai Y, Rosser CJ. Diagnostic potential of urinary α1-antitrypsin and apolipoprotein E in the detection of bladder cancer. J Urol 2012;188:2377-83.
- Tilki D, Burger M, Dalbagni G, Grossman HB, Hakenberg OW, Palou J, et al. Urine markers for detection and surveillance of non-muscle-invasive bladder cancer. Eur Urol 2011;60:484-92.
- 23. Pardoll DM, Vogelstein B, Coffey DS. A fixed site of DNA replication in eucaryotic cells. Cell 1980;19:527-36.
- 24. Berezney R, Coffey DS. Identification of a nuclear protein matrix. Biochem Biophys Res Commun 1974;60:1410-7.

- Laudadio J, Keane TE, Reeves HM, Savage SJ, Hoda RS, Lage JM, et al. Fluorescence in situ hybridization for detecting transitional cell carcinoma: implications for clinical practice. BJU Int 2005;96:1280-5.
- Glas AS, Roos D, Deutekom M, Zwinderman AH, Bossuyt PM, Kurth KH. Tumor markers in the diagnosis of primary bladder cancer. A systematic review. J Urol 2003;169:1975-82.
- Hatzichristodoulou G, Kübler H, Schwaibold H, Wagenpfeil S, Eibauer C, Hofer C, et al. Nuclear matrix protein 22 for bladder cancer detection: comparative analysis of the BladderChek*and ELISA. Anticancer Res 2012;32:5093-7.
- Wang Z, Que H, Suo C, Han Z, Tao J, Huang Z, et al. Evaluation of the NMP22 BladderChek test for detecting bladder cancer: a systematic review and meta-analysis. Oncotarget 2017;8:100648-56.
- Doğan C, Pelit ES, Yıldırım A, Zemheri IE, Çanakcı C, Başok EK, et al. The value of the NMP22 test for superficial bladder cancer diagnosis and follow-up. Turk J Urol 2013;39:137-42.
- Ponsky LE, Sharma S, Pandrangi L, Kedia S, Nelson D, Agarwal A, et al. Screening and monitoring for bladder cancer: refining the use of NMP22. J Urol 2001;166:75-8.
- Chou R, Gore JL, Buckley D, Fu R, Gustafson K, Griffin JC, et al. Urinary biomarkers for diagnosis of bladder cancer: a systematic review and meta-analysis. Ann Intern Med 2015;163:922-31.
- Sarosdy MF, Hudson MA, Ellis WJ, Soloway MS, deVere White R, Sheinfeld J, et al. Improved detection of recurrent bladder cancer using the Bard BTA stat Test. Urology 1997;50:349-53.
- 33. Heicappell R, Müller M, Fimmers R, Miller K. Qualitative determination of urinary human complement factor H-related protein (hcfHrp) in patients with bladder cancer, healthy controls, and patients with benign urologic disease. Urol Int 2000;65:181-4.
- Pode D, Shapiro A, Wald M, Nativ O, Laufer M, Kaver I. Noninvasive detection of bladder cancer with the BTA stat test. J Urol 1999;161:443-6.
- 35. Ellis WJ, Blumenstein BA, Ishak LM, Enfield DL. Clinical evaluation of the BTA TRAK assay and comparison to voided urine cytology and the Bard BTA test in patients with recurrent bladder tumors. The Multi Center Study Group. Urology 1997;50:882-7.
- 36. Thomas L, Leyh H, Marberger M, Bombardieri E, Bassi P, Pagano F, et al. Multicenter trial of the quantitative BTA TRAK assay in the detection of bladder cancer. Clin Chem 1999;45:472-7.
- 37. Comploj E, Mian C, Ambrosini-Spaltro A, Dechet C, Palermo S, Trenti E, et al. uCyt+/ImmunoCyt and cytology in the detection of urothelial carcinoma: an update on 7422

analyses. Cancer Cytopathol 2013;121:392-7.

- Hajdinjak T. UroVysion FISH test for detecting urothelial cancers: meta-analysis of diagnostic accuracy and comparison with urinary cytology testing. Urol Oncol 2008;26:646-51.
- 39. Yoder BJ, Skacel M, Hedgepeth R, Babineau D, Ulchaker JC, Liou LS, et al. Reflex UroVysion testing of bladder cancer surveillance patients with equivocal or negative urine cytology: a prospective study with focus on the natural history of anticipatory positive findings. Am J Clin Pathol 2007;127:295-301.
- Bubendorf L, Grilli B, Sauter G, Mihatsch MJ, Gasser TC, Dalquen P. Multiprobe FISH for enhanced detection of bladder cancer in voided urine specimens and bladder washings. Am J Clin Pathol 2001;116:79-86.
- 41. Sokolova IA, Halling KC, Jenkins RB, Burkhardt HM, Meyer RG, Seelig SA, et al. The development of a multitarget, multicolor fluorescence in situ hybridization assay for the detection of urothelial carcinoma in urine. J Mol Diagn 2000;2:116-23.
- 42. Savic S, Zlobec I, Thalmann GN, Engeler D, Schmauss M, Lehmann K, et al. The prognostic value of cytology and fluorescence in situ hybridization in the follow-up of nonmuscleinvasive bladder cancer after intravesical Bacillus Calmette-Guérin therapy. Int J Cancer 2009;124:2899-904.
- 43. Kamat AM, Willis DL, Dickstein RJ, Anderson R, Nogueras-González G, Katz RL, et al. Novel fluorescence in situ hybridization-based definition of bacille Calmette-Guérin (BCG) failure for use in enhancing recruitment into clinical trials of intravesical therapies. BJU Int 2016;117:754-60.
- 44. Mian C, Lodde M, Comploj E, Palermo S, Mian M, Maier K, et al. The value of the ImmunoCyt/uCyt+ test in the detection and follow-up of carcinoma in situ of the urinary bladder. Anticancer Res 2005;25:3641-4.
- 45. Lodde M, Mian C, Negri G, Berner L, Maffei N, Lusuardi L, et al. Role of uCyt+ in the detection and surveillance of urothelial carcinoma. Urology 2003;61:243-7.
- 46. Pfister C, Chautard D, Devonec M, Perrin P, Chopin D, Rischmann P, et al. Immunocyt test improves the diagnostic accuracy of urinary cytology: results of a French multicenter study. J Urol 2003;169:921-4.
- Mian C, Pycha A, Wiener H, Haitel A, Lodde M, Marberger M. Immunocyt: a new tool for detecting transitional cell cancer of the urinary tract. J Urol 1999;161:1486-9.
- Toma MI, Friedrich MG, Hautmann SH, Jäkel KT, Erbersdobler A, Hellstern A, et al. Comparison of the ImmunoCyt test and urinary cytology with other urine tests in the detection and surveillance of bladder cancer. World J Urol 2004;22:145-9.
- 49. Têtu B, Tiguert R, Harel F, Fradet Y. ImmunoCyt/uCyt+™ im-

proves the sensitivity of urine cytology in patients followed for urothelial carcinoma. Mod Pathol 2005;18:83-9.

- 50. Fradet Y, Lockhard C. Performance characteristics of a new monoclonal antibody test for bladder cancer: ImmunoCyt trade mark. Can J Urol 1997;4:400-5.
- He H, Han C, Hao L, Zang G. ImmunoCyt test compared to cytology in the diagnosis of bladder cancer: a meta-analysis. Oncol Lett 2016;12:83-8.
- Lokeshwar VB, Schroeder GL, Selzer MG, Hautmann SH, Posey JT, Duncan RC, et al. Bladder tumor markers for monitoring recurrence and screening comparison of hyaluronic acid-hyaluronidase and BTA-Stat tests. Cancer 2002;95:61-72.
- 53. Lokeshwar VB, Habuchi T, Grossman HB, Murphy WM, Hautmann SH, Hemstreet GP 3rd, et al. Bladder tumor markers beyond cytology: International Consensus Panel on bladder tumor markers. Urology 2005;66(6 Suppl 1):35-63.
- 54. Heller G, Babinsky VN, Ziegler B, Weinzierl M, Noll C, Altenberger C, et al. Genome-wide CpG island methylation analyses in non-small cell lung cancer patients. Carcinogenesis 2013;34:513-21.
- 55. Kim JG, Takeshima H, Niwa T, Rehnberg E, Shigematsu Y, Yoda Y, et al. Comprehensive DNA methylation and extensive mutation analyses reveal an association between the CpG island methylator phenotype and oncogenic mutations in gastric cancers. Cancer Lett 2013;330:33-40.
- 56. Ying J, Li H, Seng TJ, Langford C, Srivastava G, Tsao SW, et al. Functional epigenetics identifies a protocadherin PCDH10 as a candidate tumor suppressor for nasopharyngeal, esophageal and multiple other carcinomas with frequent methylation. Oncogene 2006;25:1070-80.
- Martinez VG, Munera-Maravilla E, Bernardini A, Rubio C, Suarez-Cabrera C, Segovia C, et al. Epigenetics of bladder cancer: where biomarkers and therapeutic targets meet. Front Genet 2019;10:1125.
- Zhu S, Yu W, Yang X, Wu C, Cheng F. Traditional classification and novel subtyping systems for bladder cancer. Front Oncol 2020;10:102.
- Luo Q, Vögeli TA. A methylation-based reclassification of bladder cancer based on immune cell genes. Cancers (Basel) 2020;12:3054.
- 60. Feinberg AP. Phenotypic plasticity and the epigenetics of human disease. Nature 2007;447:433-40.
- 61. Laird PW. The power and the promise of DNA methylation markers. Nat Rev Cancer 2003;3:253-66.
- 62. Witjes JA, Morote J, Cornel EB, Gakis G, van Valenberg FJP, Lozano F, et al. Performance of the Bladder EpiCheck[™] methylation test for patients under surveillance for non-muscleinvasive bladder cancer: results of a multicenter, prospective,

blinded clinical trial. Eur Urol Oncol 2018;1:307-13.

- 63. Wasserstrom A, Frumkin D, Dotan Z, Bukin E, Gadish T, Hanuka S, et al. Mp13-15 Molecular urine cytology – Bladder EpiCheck is a novel molecular diagnostic tool for monitoring of bladder cancer patients. J Urol 2016;195(4 Suppl):e140.
- 64. D'Andrea D, Soria F, Zehetmayer S, Gust KM, Korn S, Witjes JA, et al. Diagnostic accuracy, clinical utility and influence on decision-making of a methylation urine biomarker test in the surveillance of non-muscle-invasive bladder cancer. BJU Int 2019;123:959-67.
- 65. Trenti E, D'Elia C, Mian C, Schwienbacher C, Hanspeter E, Pycha A, et al. Diagnostic predictive value of the Bladder EpiCheck test in the follow-up of patients with non-muscleinvasive bladder cancer. Cancer Cytopathol 2019;127:465-9.
- 66. Zuiverloon TC, Beukers W, van der Keur KA, Munoz JR, Bangma CH, Lingsma HF, et al. A methylation assay for the detection of non-muscle-invasive bladder cancer (NMIBC) recurrences in voided urine. BJU Int 2012;109:941-8.
- 67. Su SF, de Castro Abreu AL, Chihara Y, Tsai Y, Andreu-Vieyra C, Daneshmand S, et al. A panel of three markers hyper- and hypomethylated in urine sediments accurately predicts bladder cancer recurrence. Clin Cancer Res 2014;20:1978-89.
- 68. Renard I, Joniau S, van Cleynenbreugel B, Collette C, NaôméC, Vlassenbroeck I, et al. Identification and validation of the methylated TWIST1 and NID2 genes through real-time methylation-specific polymerase chain reaction assays for the noninvasive detection of primary bladder cancer in urine samples. Eur Urol 2010;58:96-104.
- Abern MR, Owusu R, Inman BA. Clinical performance and utility of a DNA methylation urine test for bladder cancer. Urol Oncol 2014;32:51.e21-6.
- Fantony JJ, Abern MR, Gopalakrishna A, Owusu R, Jack Tay K, Lance RS, et al. Multi-institutional external validation of urinary TWIST1 and NID2 methylation as a diagnostic test for bladder cancer. Urol Oncol 2015;33:387.e1-6.
- 71. van Oers JM, Lurkin I, van Exsel AJ, Nijsen Y, van Rhijn BW, van der Aa MN, et al. A simple and fast method for the simultaneous detection of nine fibroblast growth factor receptor 3 mutations in bladder cancer and voided urine. Clin Cancer Res 2005;11:7743-8.
- 72. Descotes F, Kara N, Decaussin-Petrucci M, Piaton E, Geiguer F, Rodriguez-Lafrasse C, et al. Non-invasive prediction of recurrence in bladder cancer by detecting somatic TERT promoter mutations in urine. Br J Cancer 2017;117:583-7.
- 73. Pietzak EJ, Bagrodia A, Cha EK, Drill EN, Iyer G, Isharwal S, et al. Next-generation sequencing of nonmuscle invasive bladder cancer reveals potential biomarkers and rational therapeutic targets. Eur Urol 2017;72:952-9.
- 74. Allory Y, Beukers W, Sagrera A, Flández M, Marqués M,

ICUROLOGY

Márquez M, et al. Telomerase reverse transcriptase promoter mutations in bladder cancer: high frequency across stages, detection in urine, and lack of association with outcome. Eur Urol 2014;65:360-6.

- 75. Sampaio C, Batista R, Peralta P, Conceição P, Sismeiro A, Prazeres H, et al. Uromonitor[®]as a novel sensitive and specific urine-based test for recurrence surveillance of patients with non-muscle invasive bladder cancer. BioRxiv. 410738 [Preprint]. 2018 [cited 2021 Feb 26]. Available from: https://doi. org/10.1101/410738.
- 76. Batista R, Vinagre J, Prazeres H, Sampaio C, Peralta P, Conceição P, et al. Validation of a novel, sensitive, and specific urine-based test for recurrence surveillance of patients with non-muscle-invasive bladder cancer in a comprehensive multicenter study. Front Genet 2019;10:1237.
- 77. Sieverink CA, Batista RPM, Prazeres HJM, Vinagre J, Sampaio C, Leão RR, et al. Clinical validation of a urine test (Uromonitor-V2[®]) for the surveillance of non-muscle-invasive bladder cancer patients. Diagnostics (Basel) 2020;10:745.
- 78. Springer SU, Chen CH, Rodriguez Pena MDC, Li L, Douville C, Wang Y, et al. Non-invasive detection of urothelial cancer through the analysis of driver gene mutations and aneuploidy. Elife 2018;7:e32143.
- Rodriguez Pena MDC, Springer SU, Taheri D, Li L, Tregnago AC, Eich ML, et al. Performance of novel non-invasive urine assay UroSEEK in cohorts of equivocal urine cytology. Virchows Arch 2020;476:423-9.
- 80. Beukers W, van der Keur KA, Kandimalla R, Vergouwe Y, Steyerberg EW, Boormans JL, et al. FGFR3, TERT and OTX1 as a urinary biomarker combination for surveillance of patients with bladder cancer in a large prospective multicenter study. J Urol 2017;197:1410-8.
- Kandimalla R, Masius R, Beukers W, Bangma CH, Orntoft TF, Dyrskjot L, et al. A 3-plex methylation assay combined with the FGFR3 mutation assay sensitively detects recurrent bladder cancer in voided urine. Clin Cancer Res 2013;19:4760-9.
- 82. Roperch JP, Grandchamp B, Desgrandchamps F, Mongiat-Artus P, Ravery V, Ouzaid I, et al. Promoter hypermethylation of HS3ST2, SEPTIN9 and SLIT2 combined with FGFR3 mutations as a sensitive/specific urinary assay for diagnosis and surveillance in patients with low or high-risk non-muscleinvasive bladder cancer. BMC Cancer 2016;16:704.
- van Kessel KE, Van Neste L, Lurkin I, Zwarthoff EC, Van Criekinge W. Evaluation of an epigenetic profile for the detection of bladder cancer in patients with hematuria. J Urol 2016;195:601-7.
- 84. van Kessel KE, Beukers W, Lurkin I, Ziel-van der Made A, van der Keur KA, Boormans JL, et al. Validation of a DNA

methylation-mutation urine assay to select patients with hematuria for cystoscopy. J Urol 2017;197(3 Pt 1):590-5.

- Lotan Y, O'Sullivan P, Raman JD, Shariat SF, Kavalieris L, Frampton C, et al. Clinical comparison of noninvasive urine tests for ruling out recurrent urothelial carcinoma. Urol Oncol 2017;35:531.e15-22.
- Kavalieris L, O'Sullivan P, Frampton C, Guilford P, Darling D, Jacobson E, et al. Performance characteristics of a multigene urine biomarker test for monitoring for recurrent urothelial carcinoma in a multicenter study. J Urol 2017;197:1419-26.
- 87. Koya M, Osborne S, ChemasléC, Porten S, Schuckman A, Kennedy-Smith A. An evaluation of the real world use and clinical utility of the Cxbladder Monitor assay in the followup of patients previously treated for bladder cancer. BMC Urol 2020;20:12.
- Wallace E, Higuchi R, Satya M, McCann L, Sin MLY, Bridge JA, et al. Development of a 90-minute integrated noninvasive urinary assay for bladder cancer detection. J Urol 2018;199:655-62.
- Pichler R, Fritz J, Tulchiner G, Klinglmair G, Soleiman A, Horninger W, et al. Increased accuracy of a novel mRNAbased urine test for bladder cancer surveillance. BJU Int 2018;121:29-37.
- Valenberg FJPV, Hiar AM, Wallace E, Bridge JA, Mayne DJ, Beqaj S, et al. Prospective validation of an mRNA-based urine test for surveillance of patients with bladder cancer. Eur Urol 2019;75:853-60.
- 91. D'Elia C, Pycha A, Folchini DM, Mian C, Hanspeter E, Schwienbacher C, et al. Diagnostic predictive value of Xpert Bladder Cancer Monitor in the follow-up of patients affected by non-muscle invasive bladder cancer. J Clin Pathol 2019;72:140-4.
- 92. Hurle R, Casale P, Saita A, Colombo P, Elefante GM, Lughezzani G, et al. Clinical performance of Xpert Bladder Cancer (BC) Monitor, a mRNA-based urine test, in active surveillance (AS) patients with recurrent non-muscle-invasive bladder cancer (NMIBC): results from the Bladder Cancer Italian Active Surveillance (BIAS) project. World J Urol 2020;38:2215-20.
- Jansson MD, Lund AH. MicroRNA and cancer. Mol Oncol 2012;6:590-610.
- 94. Long JD, Sullivan TB, Humphrey J, Logvinenko T, Summerhayes KA, Kozinn S, et al. A non-invasive miRNA based assay to detect bladder cancer in cell-free urine. Am J Transl Res 2015;7:2500-9.
- 95. Zhang DZ, Lau KM, Chan ES, Wang G, Szeto CC, Wong K, et al. Cell-free urinary microRNA-99a and microRNA-125b are diagnostic markers for the non-invasive screening of bladder cancer. PLoS One 2014;9:e100793.

Piao et al

- Kim SM, Kang HW, Kim WT, Kim YJ, Yun SJ, Lee SC, et al. Cell-free microRNA-214 from urine as a biomarker for nonmuscle-invasive bladder cancer. Korean J Urol 2013;54:791-6.
- 97. Sapre N, Macintyre G, Clarkson M, Naeem H, Cmero M, Kowalczyk A, et al. A urinary microRNA signature can predict the presence of bladder urothelial carcinoma in patients undergoing surveillance. Br J Cancer 2016;114:454-62.
- 98. Rivas A, Burzio V, Landerer E, Borgna V, Gatica S, Ávila R, et al. Determination of the differential expression of mitochondrial long non-coding RNAs as a noninvasive diagnosis of bladder cancer. BMC Urol 2012;12:37.
- 99. Derrien T, Johnson R, Bussotti G, Tanzer A, Djebali S, Tilgner H, et al. The GENCODE v7 catalog of human long noncoding RNAs: analysis of their gene structure, evolution, and expression. Genome Res 2012;22:1775-89.
- 100. Khalil AM, Guttman M, Huarte M, Garber M, Raj A, Rivea Morales D, et al. Many human large intergenic noncoding RNAs associate with chromatin-modifying complexes and affect gene expression. Proc Natl Acad Sci U S A 2009;106:11667-72.
- 101. Wang L, Fu D, Qiu Y, Xing X, Xu F, Han C, et al. Genomewide screening and identification of long noncoding RNAs and their interaction with protein coding RNAs in bladder urothelial cell carcinoma. Cancer Lett 2014;349:77-86.
- 102. Zhang Z, Hao H, Zhang CJ, Yang XY, He Q, Lin J. [Evaluation of novel gene UCA1 as a tumor biomarker for the detection of bladder cancer]. Zhonghua Yi Xue Za Zhi 2012;92:384-7. Chinese.
- 103. Eissa S, Matboli M, Essawy NO, Shehta M, Kotb YM. Rapid detection of urinary long non-coding RNA urothelial carcinoma associated one using a PCR-free nanoparticle-based assay. Biomarkers 2015;20:212-7.
- 104. Martínez-Fernández M, Feber A, Dueñas M, Segovia C, Rubio C, Fernandez M, et al. Analysis of the Polycomb-related lncRNAs HOTAIR and ANRIL in bladder cancer. Clin Epigenetics 2015;7:109.
- 105. Gielchinsky I, Gilon M, Abu-Lail R, Matouk I, Hochberg A, Gofrit ON, et al. H19 non-coding RNA in urine cells detects urothelial carcinoma: a pilot study. Biomarkers 2017;22:661-6.
- 106. Martens-Uzunova ES, Böttcher R, Croce CM, Jenster G, Visakorpi T, Calin GA. Long noncoding RNA in prostate, bladder, and kidney cancer. Eur Urol 2014;65:1140-51.
- 107. Quan J, Pan X, Zhao L, Li Z, Dai K, Yan F, et al. LncRNA as a diagnostic and prognostic biomarker in bladder cancer: a systematic review and meta-analysis. Onco Targets Ther 2018;11:6415-24.
- 108. Cao Y, Tian T, Li W, Xu H, Zhan C, Wu X, et al. Long noncoding RNA in bladder cancer. Clin Chim Acta 2020;503:113-

21.

- 109. Fujita K, Pavlovich CP, Netto GJ, Konishi Y, Isaacs WB, Ali S, et al. Specific detection of prostate cancer cells in urine by multiplex immunofluorescence cytology. Hum Pathol 2009;40:924-33.
- Tyler KL, Selvaggi SM. Morphologic features of prostatic adenocarcinoma on ThinPrep[®]urinary cytology. Diagn Cytopathol 2011;39:101-4.
- 111. Saetun P, Semangoen T, Thongboonkerd V. Characterizations of urinary sediments precipitated after freezing and their effects on urinary protein and chemical analyses. Am J Physiol Renal Physiol 2009;296:F1346-54.
- 112. Brinkman JA, Rahmani MZ, Jones WE, Chaturvedi AK, Hagensee ME. Optimization of PCR based detection of human papillomavirus DNA from urine specimens. J Clin Virol 2004;29:230-40.
- 113. Lin SY, Linehan JA, Wilson TG, Hoon DSB. Emerging utility of urinary cell-free nucleic acid biomarkers for prostate, bladder, and renal cancers. Eur Urol Focus 2017;3:265-72.
- 114. Yu X, Wang R, Han C, Wang Z, Jin X. A panel of urinary long non-coding RNAs differentiate bladder cancer from urocystitis. J Cancer 2020;11:781-7.
- 115. Soria F, Droller MJ, Lotan Y, Gontero P, D'Andrea D, Gust KM, et al. An up-to-date catalog of available urinary biomarkers for the surveillance of non-muscle invasive bladder cancer. World J Urol 2018;36:1981-95.
- 116. Pichler R, Tulchiner G, Fritz J, Schaefer G, Horninger W, Heidegger I. Urinary UBC Rapid and NMP22 test for bladder cancer surveillance in comparison to urinary cytology: results from a prospective single-center study. Int J Med Sci 2017;14:811-9.
- 117. Ecke TH, Weiß S, Stephan C, Hallmann S, Arndt C, Barski D, et al. UBC**Rapid* Test-a urinary point-of-care (POC) assay for diagnosis of bladder cancer with a focus on non-muscle invasive high-grade tumors: results of a multicenter-study. Int J Mol Sci 2018;19:3841.
- 118. Babjuk M, Soukup V, Pešl M, KostírováM, DrncováE, SmolováH, et al. Urinary cytology and quantitative BTA and UBC tests in surveillance of patients with pTapT1 bladder urothelial carcinoma. Urology 2008;71:718-22.
- 119. Mungan NA, Vriesema JL, Thomas CM, Kiemeney LA, Witjes JA. Urinary bladder cancer test: a new urinary tumor marker in the follow-up of superficial bladder cancer. Urology 2000;56:787-92.
- Fenner A. ADXBLADDER test better than cytology for NMIBC follow-up monitoring. Nat Rev Urol 2020;17:486.
- 121. Gontero P, Montanari E, Roupret M, Longo F, Stockley J, Kennedy A, et al. Comparison of the performances of the ADXBLADDER test and urinary cytology in the follow-up of

non-muscle-invasive bladder cancer: a blinded prospective multicentric study. BJU Int 2021;127:198-204.

- 122. Guo XG, Long JJ. Cytokeratin-19 fragment in the diagnosis of bladder carcinoma. Tumour Biol 2016;37:14329-30.
- 123. Kuang LI, Song WJ, Qing HM, Yan S, Song FL. CYFRA21-1 levels could be a biomarker for bladder cancer: a meta-analysis. Genet Mol Res 2015;14:3921-31.
- 124. Nisman B, Barak V, Shapiro A, Golijanin D, Peretz T, Pode D. Evaluation of urine CYFRA 21-1 for the detection of primary and recurrent bladder carcinoma. Cancer 2002;94:2914-22.
- 125. Bensalah K, Montorsi F, Shariat SF. Challenges of cancer biomarker profiling. Eur Urol 2007;52:1601-9.
- 126. Goebell PJ, Groshen SL, Schmitz-Dräger BJ. Guidelines for development of diagnostic markers in bladder cancer. World J Urol 2008;26:5-11.
- 127. Shariat SF, Lotan Y, Vickers A, Karakiewicz PI, Schmitz-Dräger BJ, Goebell PJ, et al. Statistical consideration for clinical biomarker research in bladder cancer. Urol Oncol 2010;28:389-400.
- 128. Cosma CL, Sherman DR, Ramakrishnan L. The secret lives of the pathogenic mycobacteria. Annu Rev Microbiol 2003;57:641-76.
- 129. Curtiss N, Balachandran A, Krska L, Peppiatt-Wildman C, Wildman S, Duckett J. Age, menopausal status and the bladder microbiome. Eur J Obstet Gynecol Reprod Biol 2018;228:126-9.
- 130. Lewis DA, Brown R, Williams J, White P, Jacobson SK, Marchesi JR, et al. The human urinary microbiome; bacterial DNA in voided urine of asymptomatic adults. Front Cell Infect Microbiol 2013;3:41.
- 131. Hilt EE, McKinley K, Pearce MM, Rosenfeld AB, Zilliox MJ, Mueller ER, et al. Urine is not sterile: use of enhanced urine culture techniques to detect resident bacterial flora in the adult female bladder. J Clin Microbiol 2014;52:871-6.
- 132. Cimadamore A, Santoni M, Massari F, Gasparrini S, Cheng L, Lopez-Beltran A, et al. Microbiome and cancers, with focus

on genitourinary tumors. Front Oncol 2019;9:178.

- 133. Akram A, Maley M, Gosbell I, Nguyen T, Chavada R. Utility of 16S rRNA PCR performed on clinical specimens in patient management. Int J Infect Dis 2017;57:144-9.
- 134. Yu Y, Pieper R. Urinary pellet sample preparation for shotgun proteomic analysis of microbial infection and host-pathogen Interactions. In: Posch A. Proteomic profiling: methods and protocols. New York: Humana Press; 2015;65-74.
- 135. Yu Y, Sikorski P, Bowman-Gholston C, Cacciabeve N, Nelson KE, Pieper R. Diagnosing inflammation and infection in the urinary system via proteomics. J Transl Med 2015;13:111.
- 136. Redelman-Sidi G, Glickman MS, Bochner BH. The mechanism of action of BCG therapy for bladder cancer--a current perspective. Nat Rev Urol 2014;11:153-62.
- 137. Xu W, Yang L, Lee P, Huang WC, Nossa C, Ma Y, et al. Minireview: perspective of the microbiome in the pathogenesis of urothelial carcinoma. Am J Clin Exp Urol 2014;2:57-61.
- 138. BučevićPopovićV, Šitum M, Chow CT, Chan LS, Roje B, TerzićJ. The urinary microbiome associated with bladder cancer. Sci Rep 2018;8:12157.
- 139. Wu P, Zhang G, Zhao J, Chen J, Chen Y, Huang W, et al. Profiling the urinary microbiota in male patients with bladder cancer in China. Front Cell Infect Microbiol 2018;8:167.
- 140. Chipollini J, Wright JR, Nwanosike H, Kepler CY, Batai K, Lee BR, et al. Characterization of urinary microbiome in patients with bladder cancer: results from a single-institution, feasibility study. Urol Oncol 2020;38:615-21.
- 141. Wolfe AJ, Toh E, Shibata N, Rong R, Kenton K, Fitzgerald M, et al. Evidence of uncultivated bacteria in the adult female bladder. J Clin Microbiol 2012;50:1376-83.
- 142. Bajic P, Van Kuiken ME, Burge BK, Kirshenbaum EJ, Joyce CJ, Wolfe AJ, et al. Male bladder microbiome relates to lower urinary tract symptoms. Eur Urol Focus 2020;6:376-82.
- 143. Jo JH, Kennedy EA, Kong HH. Research techniques made simple: bacterial 16S ribosomal RNA gene sequencing in cutaneous research. J Invest Dermatol 2016;136:e23-7.