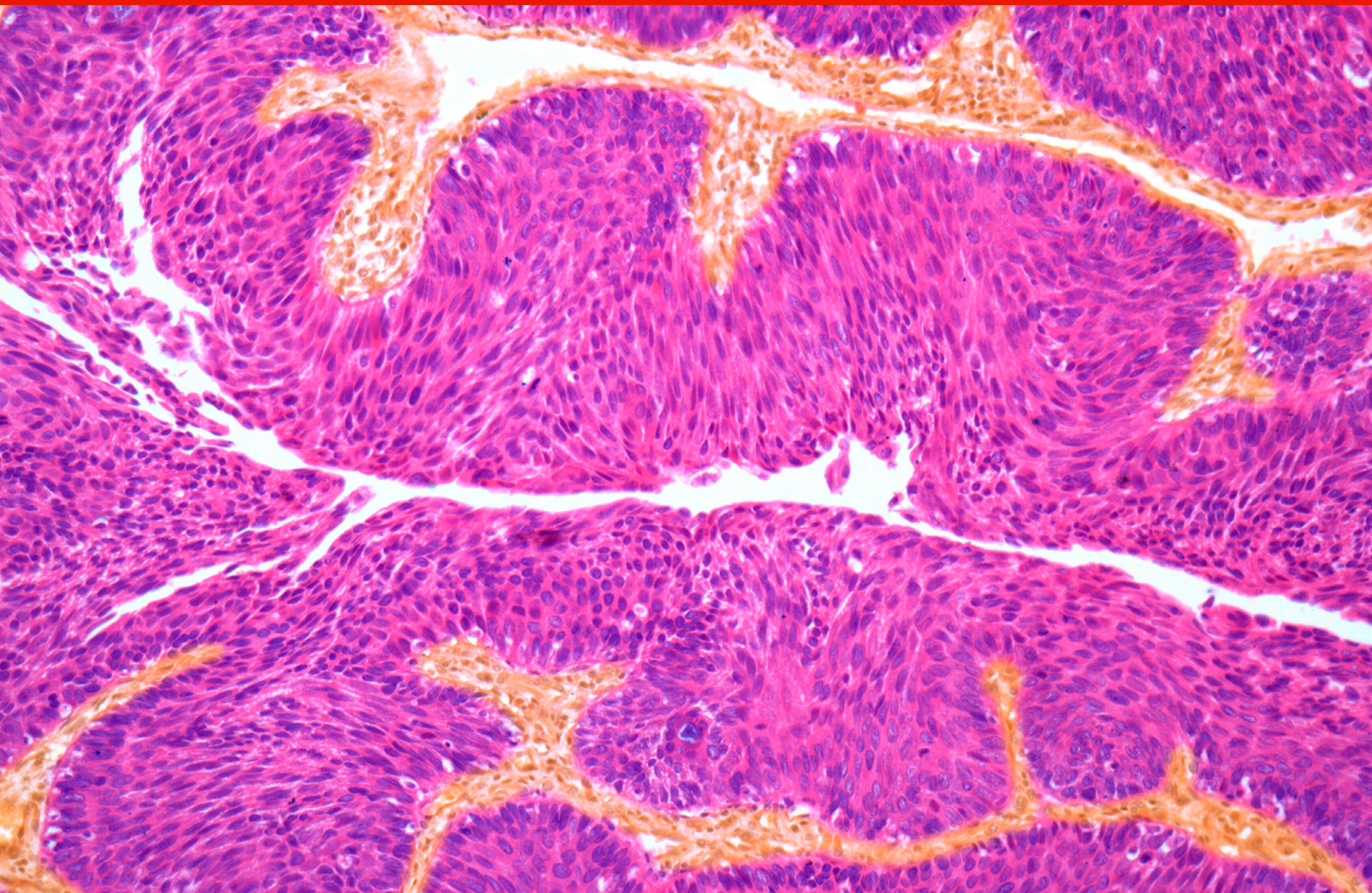


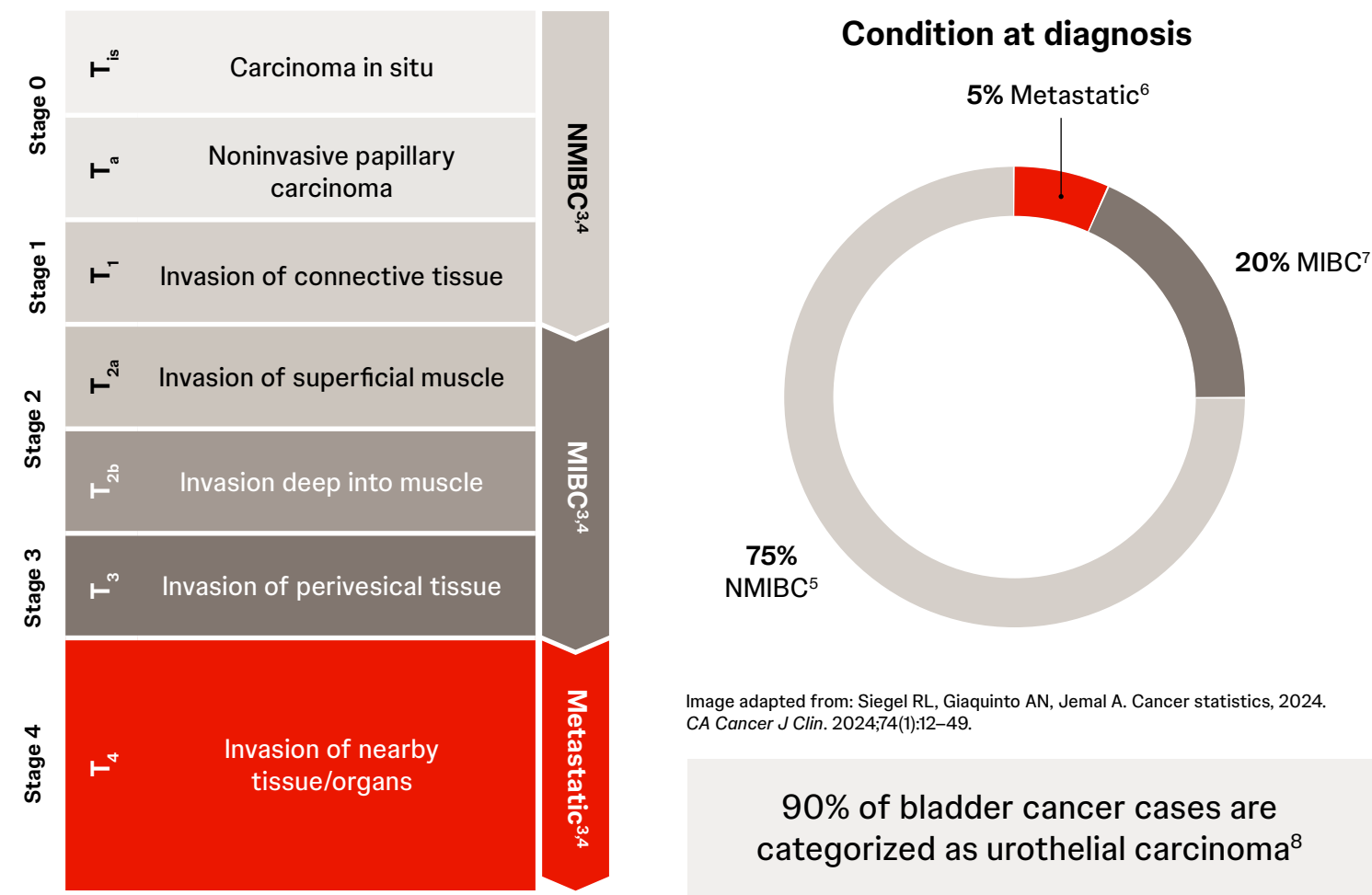
# Bladder Cancer Biomarkers and Guideline Recommendations

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Precision Medicine

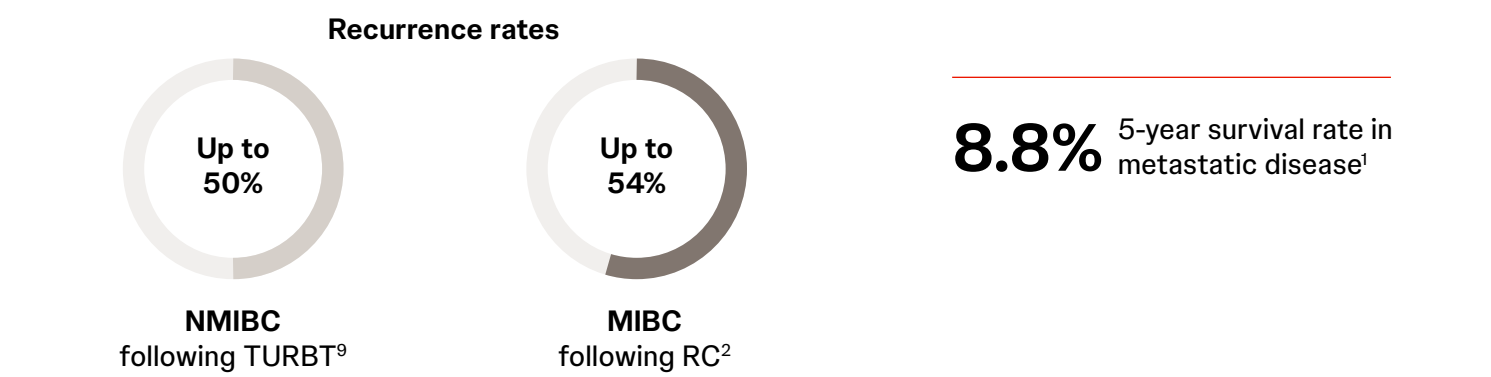


Bladder cancer is a prevalent, commonly recurring disease<sup>1,2</sup>

Bladder cancer is the sixth most prevalent cancer in the United States<sup>1</sup>

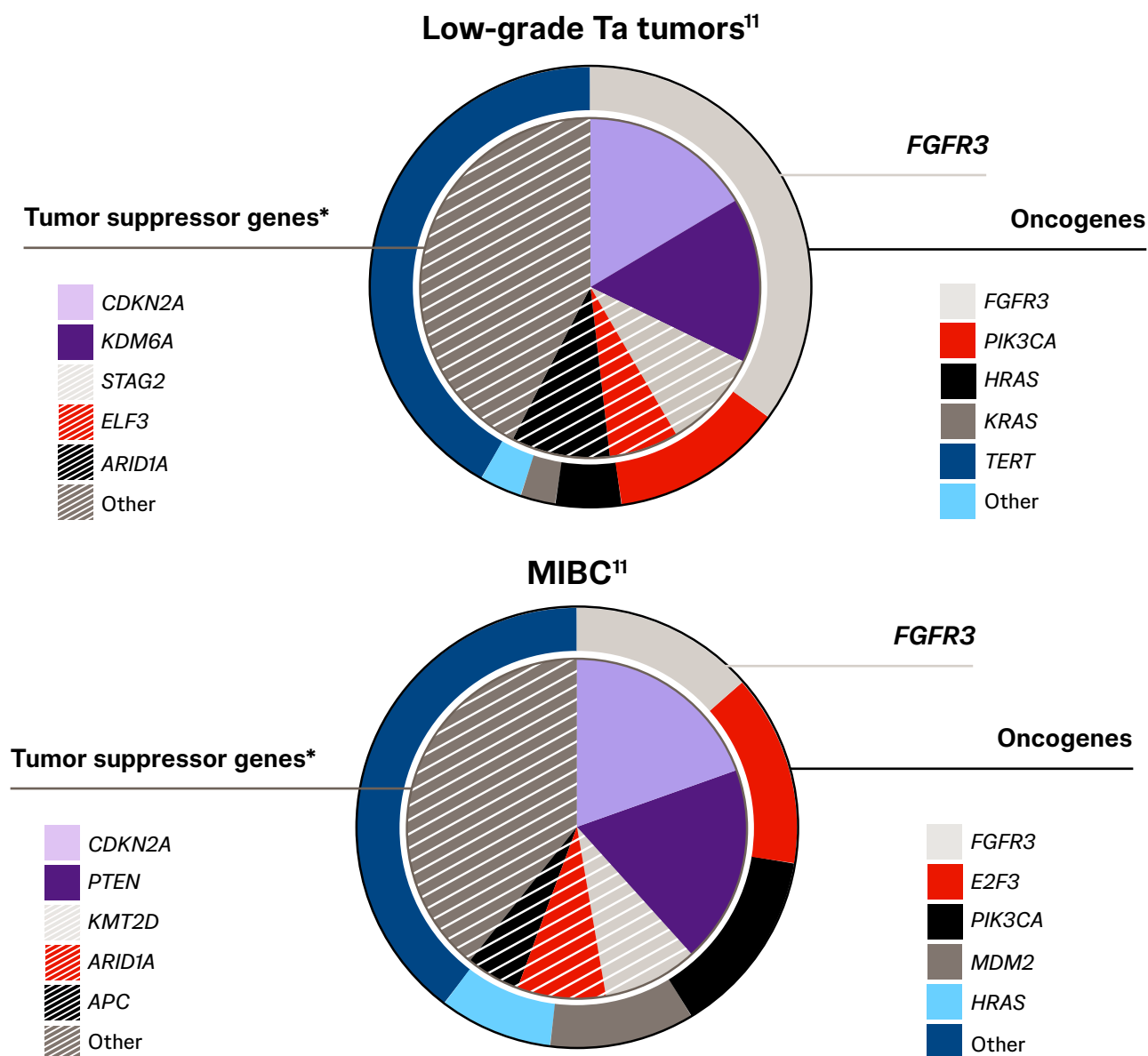


Although bladder cancer is often caught in early stages, **recurrence rates are high and survival rates in metastatic disease are low**<sup>1,2,5</sup>



## The understanding and approach to bladder cancer is evolving with precision medicine<sup>10</sup>

Urothelial carcinoma is driven by a variety of stage-dependent biomarkers<sup>11</sup>



Images adapted from: Sanli O, Dobruch J, Knowles MA, et al. Bladder cancer. *Nat Rev Dis Primers*. 2017;3:17022. doi:10.1038/nrdp.2017.22

### Opportunity to optimize

Biomarker testing may identify patients at risk of recurrence or progression and help guide treatment decisions<sup>10</sup>

\*Summed percentages exceed 100% since patients can express more than 1 mutation.



# Bladder Cancer Biomarkers and Guideline Recommendations

Biomarkers are emerging as useful diagnostic, prognostic, and predictive tools in the management of bladder cancer<sup>12-14\*</sup>

Biomarker testing at different stages of disease can inform the management of bladder cancer<sup>12,13\*</sup>



**Diagnosis**  
(eg, NMP22)<sup>14</sup>



**Understanding genetic disposition**<sup>12,15\*</sup>  
(eg, MSH2)<sup>15,16\*</sup>



**Risk stratification and prognosis**  
(eg, CDKN2A)<sup>17</sup>



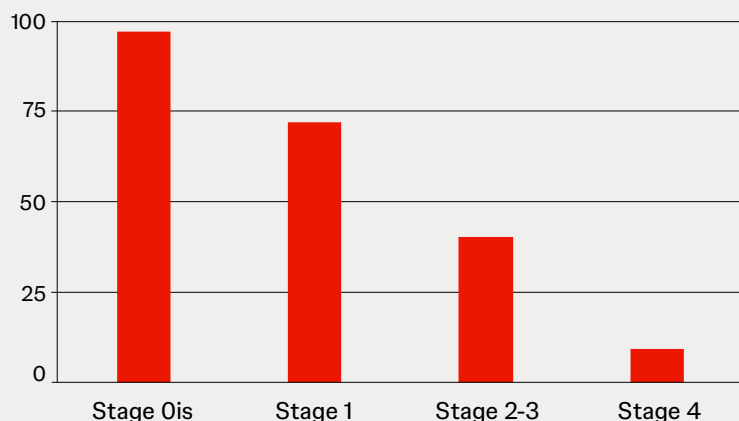
**Predict likelihood of treatment response**  
(eg, FGFR3)<sup>12\*</sup>



**Screen for clinical trials**<sup>12\*</sup>  
(eg, ARID1A)<sup>18</sup>

Early diagnosis is associated with better clinical outcomes<sup>1</sup>

**Five-year relative survival rates**<sup>1,19</sup>



Survival rates are correlated with cancer stage at detection<sup>1</sup>

Biomarker analysis from liquid biopsies are emerging as potential noninvasive diagnostic tools<sup>20</sup>

Biomarker testing can significantly improve accuracy and overcome current limitations in bladder cancer evaluation<sup>20-22</sup>

- Testing may aid in interpreting equivocal results from cytology<sup>13</sup>
- Repeat testing may help monitor patients’ response to therapy<sup>13,22,23</sup>
- Matching patients with bladder cancer to biomarker-informed treatments can lead to clinical benefits<sup>24-27</sup>

Clinical guidelines for bladder cancer recommend biomarker testing at diagnosis and when monitoring treatment response<sup>12,13,28\*</sup>

When to order testing:	Following diagnosis	When monitoring treatment response
National Comprehensive Cancer Network® (NCCN®) <sup>12*</sup>	At time of diagnosis of advanced disease <ul style="list-style-type: none"><li>• Facilitate treatment selection</li><li>• Prevent delays in:<ul style="list-style-type: none"><li>- Later lines of therapy</li><li>- Entry into clinical trials</li></ul></li></ul>	
American Urological Association (AUA)/ Society of Urologic Oncology (SUO) <sup>13</sup>	Following initial diagnosis <ul style="list-style-type: none"><li>• Resolve equivocal cytology (UroVysion® FISH and ImmunoCyt)</li></ul>	<ul style="list-style-type: none"><li>• Assess treatment response to intravesical BCG in NMIBC</li></ul>
Emerging evidence:  NCCN <sup>12*</sup>		<ul style="list-style-type: none"><li>• Can be considered for the surveillance of high-risk NMIBC patients</li></ul>
AUA/American Society of Clinical Oncology (ASCO)/SUO <sup>13,28,29</sup>		<ul style="list-style-type: none"><li>• To identify early recurrences post surgery using ctDNA<ul style="list-style-type: none"><li>- Determine patients that are likely (or not likely) to benefit from adjuvant treatment</li></ul></li></ul>

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# Noninvasive biomarkers for bladder cancer detection are continuing to emerge<sup>14,28</sup>

As of 2024, there are currently **6 FDA-approved tests for bladder cancer diagnosis and monitoring based on urinary biomarkers**, including:

UroVysion®, ImmunoCyt®, BTA Stat, BTA TRAK, NMP22® BladderChek®, and NMP22® Bladder Cancer Test<sup>14,30</sup>

Recommended by AUA/SUO NMIBC guideline<sup>13</sup>

<b>UroVysion®</b> Chromosomal alterations <sup>14</sup>  <b>Chromosome 3, 7, or 17 aneuploidy and loss of the 9p21 locus<sup>14</sup></b>  Used to assess response to intravesical BCG and resolve equivocal cytology <sup>13</sup>	<b>ImmunoCyt®</b> Bladder cancer cell antigens <sup>14</sup>  Detection of cells positive for <b>carcinoembryonic antigens and sulfated mucin glycoproteins</b> may aid in bladder cancer management <sup>14</sup>  Used to resolve equivocal cytology <sup>13</sup>
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## Did you know?

According to NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines®), urinary biomarkers may be considered in the first 2 years of surveillance for high-risk NMIBC<sup>12\*</sup>

# Biomarker testing may inform the likelihood of treatment response in advanced disease<sup>12\*</sup>

Biomarker	Prevalence in locally advanced or metastatic disease	Testing methods†	Predict response to
FGFR3 alteration‡	15% <sup>31</sup>	RT-PCR, NGS, FISH <sup>32–34</sup>	FGFR inhibitors <sup>12,24*</sup>
HER2 overexpression	13% <sup>35</sup>	IHC, FISH <sup>35</sup>	HER2-targeted therapy <sup>12,35*</sup>
MSI-H	<1% <sup>27</sup>	NGS, RT-PCR <sup>27</sup>	Immune checkpoint inhibitors <sup>12,27,36*</sup>
High TMB	30% <sup>36</sup>	NGS <sup>37</sup>	
PD-L1 overexpression	35% <sup>38</sup>	IHC <sup>38</sup>	

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†ctDNA is not currently utilized in clinical care to inform treatment decisions.<sup>33,39</sup>  
‡A larger panel may be preferred to identify rare mutations that may have approved therapies or allow for clinical trial eligibility.<sup>12\*</sup>

## *FGFR3* alterations are predictive biomarkers in bladder cancer<sup>12\*</sup>

Oncogenic *FGFR* alterations, including point mutations and fusions, may cause constitutive signaling<sup>32,40</sup>

- Promotes tumorigenesis by increasing survival, migration, proliferation, angiogenesis, and invasion

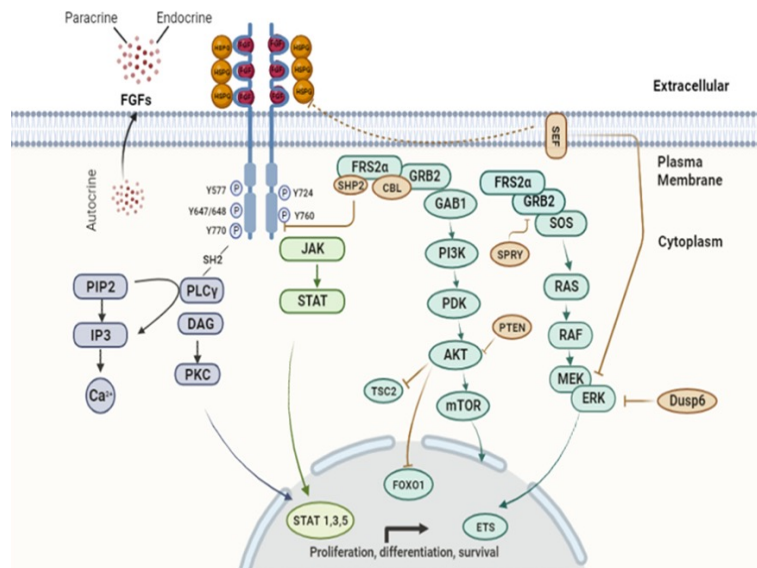
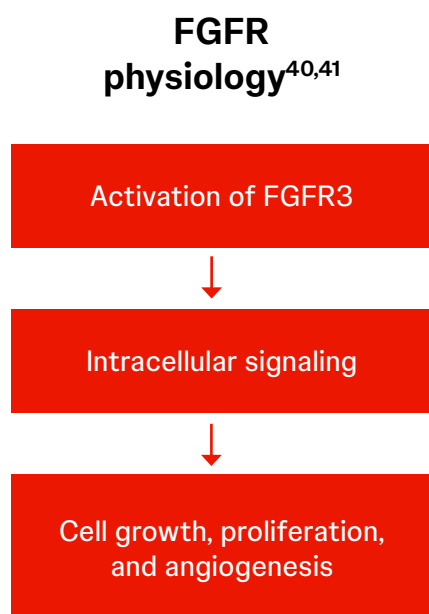


Image adapted from: Ascione CM, Napolitano F, Esposito D, et al. Role of *FGFR3* in bladder cancer: treatment landscape and future challenges. *Cancer Treat Rev.* 2023;115:102530. doi:10.1016/j.ctrv.2023.102530

### Clinically relevant *FGFR3* alterations

#### ***FGFR3* mutations:**

p.R248C, p.S249C, p.G370C, and p.Y373C<sup>42</sup>

- Frequency is stage-dependent
  - ~75% in NMIBC; ~20% in MIBC; ~9% in mMIBC<sup>43,44</sup>

#### ***FGFR3* fusions:**

*FGFR3*-TACC3v1 and *FGFR3*-TACC3v3<sup>33,43</sup>

- 2–6% in patients with MIBC or mMIBC<sup>42,43</sup>

### Did you know?

*FGFR3* alterations, including clinically relevant point mutations and fusions, can be detected with PCR, NGS, and FISH<sup>32–34</sup>

# Some urothelial tumors express predictive biomarkers for treatments with pan tumor indications<sup>12,16,45,46\*</sup>

### Predictive biomarkers (pan tumor)

	<i>HER2</i> <sup>12*</sup>	TMB <sup>47</sup>	MSI and dMMR <sup>45</sup>
Method	IHC – overexpression <sup>48</sup> FISH – amplification <sup>48</sup> NGS – overexpression and amplification <sup>49,50</sup>	NGS – whole-genome sequencing (WGS) <sup>51</sup>	PCR – microsatellite markers <sup>16,45</sup> IHC – expression of MMR proteins <sup>16,45</sup> NGS – targeted gene panels <sup>16,45</sup>
Clinical considerations	<ul style="list-style-type: none"><li>IHC 2+ scores are equivocal<sup>48</sup><ul style="list-style-type: none"><li>follow up with FISH<sup>48</sup></li></ul></li></ul>	<ul style="list-style-type: none"><li>High TMB &gt;10 mutations/MB<sup>36</sup></li></ul>	<ul style="list-style-type: none"><li>Loss of ≥1 MMR protein defines dMMR<sup>45</sup><ul style="list-style-type: none"><li>MLH1, PMS2, MSH2 and MSH6<sup>16</sup></li></ul></li><li>MSI-H: instability at &gt;30% of loci<sup>52</sup></li></ul>

### Did you know?

Although less common in bladder cancer, treatments are available for solid tumors that have metastasized and express specific alterations in *NTRK1/2/3*, *RET*, or *BRAF*<sup>53–55</sup>

Testing for biomarkers including *FGFR3*, *HER2*, MSI, dMMR, and TMB can identify patients who are eligible for biomarker-informed therapies<sup>12,45,47\*</sup>



# Emerging Biomarkers and Future Directions

New biomarkers are being identified that can potentially improve multiple domains of bladder cancer management<sup>18,56–70</sup>

Extracellular vesicles are emerging as useful sources of biomarkers<sup>56,57</sup>

	DNA <sup>18,58–60</sup> Mutations, fusions, methylation	RNA <sup>61–63</sup> mRNA, miRNA, ncRNA, lncRNA	Protein <sup>57,64–70</sup> Overexpression
Diagnosis*	FGFR3, PIK3CA, KRAS, TERT, TP53	MALAT, PCAT-1, SPRY4-IT1	alpha-1-anti-trypsin, BLCA-1/4, H2B1K
Prognosis*	CDKN2A, CTSV, FOXM1	H19, PCAT-1, UBC1	EphA2, ERCC1/2, Rab1a
Prediction of treatment response*	ARID1A, KDM6A, KMT2D		EphA2, HER2, HER3

## Opportunity to optimize

An MDT approach can help ensure that patients’ biomarker profiles are continuously updated based on current approvals<sup>71</sup>

\*This is not an exhaustive list of emerging biomarkers for bladder cancer.

# Bladder Cancer Biomarkers and Guideline Recommendations

## Summary



Bladder cancer is a molecularly heterogeneous disease; different tumor subtypes and stages express unique molecular signatures<sup>11</sup>



Multiple guidelines recommend that biomarker testing may be utilized to aid in diagnosis and prognosis or to inform treatment decisions in select bladder cancer patients<sup>12,13\*</sup>



Biomarkers may play an instrumental role in diagnosis, risk stratification, surveillance, and identifying treatment options<sup>12,13,28\*</sup>

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APC, adenomatous polyposis coli; ARID1A, AT-rich interactive domain-containing protein 1A; AUA, American Urological Association; BCG, Bacillus Calmette-Guérin; BLCA, bladder cancer; BRAF, B-Raf proto-oncogene; CDKN2A, cyclin-dependent kinase inhibitor 2A; ctDNA, circulating tumor deoxyribonucleic acid; CTSV, cathepsin V; dMMR, deficient DNA mismatch repair; DNA, deoxyribonucleic acid; E2F3, E2F transcription factor 3; ELF3, E74-like factor 3; EphA2, ephrin type-A receptor 2; ERCC, excision repair 1, endonuclease non-catalytic subunit; FDA, Food and Drug Administration; FGFR, fibroblast growth factor receptor; FISH, fluorescence in situ hybridization; FOXM1, forkhead box M1; H2B1K, histone H2B type 1-K; HER2, human epidermal growth factor receptor 2; HER3, human epidermal growth factor receptor 3; HRAS, Harvey rat sarcoma viral oncogene homologue; IHC, immunohistochemistry; KDM6A, lysine-specific demethylase 6A; KMT2D, Histone-lysine N-methyltransferase 2D; KRAS, Kirsten rat sarcoma viral oncogene homologue; lncRNA, long non-coding ribonucleic acid; MALAT, metastasis associated lung adenocarcinoma transcript; MB, megabase; MDM2, murine double minute 2; MDT, multidisciplinary team; MIBC, muscle-invasive bladder cancer; miRNA, micro ribonucleic acid; mMIBC, metastatic muscle-invasive bladder cancer; MLH1, MutL homologue 1; MMR, mismatch repair; mRNA, messenger ribonucleic acid; MSH, MutS homologue; MSI, microsatellite instability; MSI-H, microsatellite instability-high; NCCN, National Comprehensive Cancer Network; ncrRNA, non-coding ribonucleic acid; NGS, next-generation sequencing; NMIBC, non-muscle-invasive bladder cancer; NMP22, nuclear matrix protein 22; NTRK, neurotrophic tyrosine receptor kinase; PCAT-1, prostate cancer associated transcript 1; PCR, polymerase chain reaction; PD-L1, programmed cell death ligand 1; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha; PMS2, postmeiotic segregation increased 2; Rab1a, Ras-related protein Rab-1A; RC, radical cystectomy; RET, rearranged during transfection; RNA, ribonucleic acid; RT-PCR, reverse transcription polymerase chain reaction; SPRY4-T1, sprouty receptor tyrosine kinase signaling antagonist 4 transcript 1; STAG2, stromal antigen 2; SUO, Society of Urologic Oncology; TACC3, transforming acidic coiled-coil-containing protein 3; TERT, telomerase reverse transcriptase; TMB, tumor mutational burden; TP53, tumor protein p53; TURBT, transurethral resection of bladder tumor; UBC1, Ubiquitin-conjugating enzyme E2 1.

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