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## CTC-derived AR-V7 detection as a prognostic and predictive biomarker in advanced prostate cancer

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### Abstract

**Introduction:** Prostate cancer is a highly heterogeneous disease, with remarkably different prognosis across all stages. Increased circulating tumor cell (CTC) count (  $\geq 5$ ) using the CellSearch assay has been identified as one of the markers that can be used to predict survival, with added value beyond currently available prognostic factors. Recently, androgen receptor splice variant 7 (AR-V7) detection has been associated with worse outcomes for patients with castration-resistant prostate cancer (CRPC) treated with novel androgen receptor-signaling (ARS) inhibitors such as abiraterone and enzalutamide but not taxane chemotherapies.

**Areas covered:** In this manuscript, the authors review the available biomarkers in CRPC and discuss emerging data on the value of CTC-derived AR-V7 status to assess prognosis and its potential role to guide treatment selection for patients with advanced prostate cancer.

**Expert Commentary:** Current evidence supports AR-V7 status as a prognostic biomarker and also as a potential predictive biomarker for patients with mCRPC. The authors expect that the incorporation of AR-V7 status and other biomarkers (e.g. AR mutations) in the sequential assessment of patients with advanced prostate cancer will lead to a more rational use of available and future therapies, with significant improvements in outcomes for our patients.

### Keywords

AR-V7; prognostic; predictive; biomarker; castration-resistant prostate cancer

## 1. INTRODUCTION

Prostate cancer remains one of the leading causes of cancer death worldwide<sup>1,2</sup> despite many recent advances in the understanding of the disease biology and increasing treatment options for patients with advanced prostate cancer.

For many decades androgen deprivation therapy (ADT) has been the mainstay of therapy for patients with metastatic prostate cancer, with high rates of PSA declines, symptom improvement and disease control. Despite this initial favorable response, most patients

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eventually develop resistance to ADT and present with disease progression. Over the past decade we have learned that there are several different mechanisms of progression in the setting of castrate levels of testosterone, but most patients present with a rising PSA, reflecting continuous androgen receptor (AR) signaling activation<sup>3</sup>.

Better understanding of the mechanisms of disease progression has led to the development and approval of several active systemic therapies for castration-resistant prostate cancer (CRPC) over the last several years (Table 1), including novel AR-signaling inhibitors (abiraterone<sup>4,5</sup> and enzalutamide<sup>6,7</sup>), taxane chemotherapy with docetaxel<sup>8</sup> and cabazitaxel<sup>9</sup>, and agents with alternative mechanisms of action such as radium-223<sup>10</sup> and sipuleucel-T<sup>11</sup>. Recently, clinical trials evaluating the use of docetaxel and abiraterone in castration-sensitive prostate cancer (CSPC) have demonstrated further improvements in outcomes and increased survival compared to ADT alone<sup>12–15</sup>, particularly in those with metastatic castration-sensitive disease, suggesting that earlier use of active drugs can lead to even more profound clinical benefits.

Despite these recent achievements, many patients derive short-term benefits from available therapies and even those who have a prolonged disease control will still develop eventual progressive disease at some point. It is widely known that prostate cancer is a highly heterogeneous disease, with some patients presenting with very indolent disease and others with highly aggressive clinical course and refractory disease. Therefore, it is of paramount importance to establish prognosis in this setting but also to identify predictive biomarkers of benefit to specific therapies (*i.e.* treatment-selection markers) to help us select the best drug for the appropriate patient at the appropriate time and with the appropriate toxicities (*i.e.*, “precision medicine”). In this article, we will review the available prognostic markers for advanced prostate cancer, with a focus on CTC-detected AR splice variant 7 (AR-V7), and also to describe potential predictive biomarkers in this setting.

## 2. MECHANISMS OF ESCAPE TO ANDROGEN RECEPTOR-SIGNALING INHIBITORS

The mechanisms of resistance to ADT and other AR-signaling (ARS) inhibitors for metastatic prostate cancer can be broadly divided into androgen-dependent and androgen-independent groups<sup>16–18</sup> (Figure 1). Despite not being the main subject of this review, it is important to recognize that many of these resistance mechanisms have prognostic implications since they usually lead to less responsiveness to other available therapies and consequently to a worse overall prognosis. One of the most important examples of a prognostic and predictive marker is the detection of AR-V7 in circulating tumor cells (CTCs), which has been demonstrated in several single-center studies to be associated with a poor prognosis, as discussed in detail over the next several sections of this paper.

## 3. BIOMARKERS IN METASTATIC PROSTATE CANCER

### 3.1. Biomarker definitions

The term biomarker refers to clinical or molecular characteristics that can be objectively and reproducibly measured to indicate a biologic condition, including normal or pathogenic

processes, and also a response to a specific therapeutic intervention<sup>19,20</sup>. Therefore, biomarkers can be clinical features (such as performance status), laboratory analytes (such as hemoglobin, PSA, etc.), imaging studies, or molecular alterations (such as gene mutations). These biomarkers are classified based on their contexts of use (Table 2). Importantly, each biomarker under development must undergo the prerequisite analytical and clinical validation steps before it can be reliably used to inform a medical decision. Analytical validation comprises several steps to guarantee accuracy and reproducibility of the biomarker assay and the clinical validation is established when clinical trials demonstrate that the specific biomarker provides useful information to guide management<sup>19,20</sup>.

Several biomarkers have been identified as prognostic factors for patients with advanced prostate cancer<sup>21</sup>. Many of these markers that are associated with clinically meaningful outcomes, such as survival, can be identified at baseline and others can be identified after treatment over the course of the disease (Table 3). A useful prognostic model for predicting OS in patients with mCRPC treated with first-line chemotherapy has been developed using several of these markers<sup>22</sup>.

### 3.2. Circulating Tumor Cell Count and Prognosis

Circulating tumor cell enumeration in patients with mCRPC has been correlated with survival in several studies, both retrospective and prospectively<sup>23–25</sup>. It has been demonstrated that patients with a CTC count  $\geq 5$  at baseline have worse survival and also that patients with a CTC count decline (from  $\geq 5$  to  $< 5$ ) after 12 weeks of therapy have better outcomes than patients with no CTC declines<sup>26</sup>. Two recent studies have demonstrated that incorporating the CTC count with other known prognostic markers significantly improves the prognostic assessment and may serve as a surrogate for survival in mCRPC<sup>27,28</sup>. In the COU-AA-301 phase III randomized trial comparing abiraterone plus prednisone versus placebo plus prednisone for patients with mCRPC previously treated with docetaxel, the analysis of CTC count and other known prognostic biomarkers (PSA, LDH, hemoglobin [Hb], albumin [Alb] and alkaline phosphatase levels [AlkPhos]) as a surrogate for survival was prospectively assessed as a secondary endpoint. The final analysis included 711 patients and demonstrated that a panel of CTC count plus LDH level was able to stratify patients according to three different groups in terms of prognosis: low (CTC  $< 5$ ; any LDH), intermediate (CTC  $\geq 5$  and LDH  $\leq 250$ ), and high-risk (CTC  $\geq 5$  and LDH  $> 250$ ), with median OS being 8.7, 12.0 and 22.2 months, respectively ( $P < 0.001$ ), satisfying Prentice surrogacy criteria for survival<sup>27</sup>. Another recent study examined the added prognostic value of CTC count to a model containing LDH, PSA, Hb, Alb and AlkPhos in mCRPC patients enrolled in two prospective randomized trials (COU-AA-301 and ELM-PC4)<sup>28</sup>. This study also demonstrated that including the CTC count in current prognostic models increased its discriminating power for predicting survival in mCRPC patients.

It should be noted that most of these studies on CTC enumeration as a prognostic biomarker were performed using the CellSearch assay, which relies on capturing CTCs of epithelial origin from whole blood, and is the only FDA cleared method for CTC detection. There are some limitations of the CellSearch assay, including its limited sensitivity since it can only detect EPCAM+ cells (would miss cells undergoing mesenchymal transition), and the

uncertainty of whether detected cells are truly viable cancer cells rather than necrotic cells. Thus, many other assays are under development to improve CTC detection sensitivity by including other CTC detection methods, such as the use of multiple antibodies (CK, HER-2, CD44, CD24, ALDH1) and also DNA- or RNA-based RT-PCR assays. A detailed review of CTC detection methods has been reported elsewhere<sup>29</sup>.

### 3.3. AR-V7 as a Prognostic and Predictive Biomarker in CRPC

Androgen receptor splice variants are constitutively active forms of the AR that lack the ligand-binding domain and have been implicated as one of the mechanisms of resistance to the currently available ARS inhibitors. Most of these variants arise from abnormal splicing (and retention) of intronic sequences containing cryptic exons which (when translated) result in frameshift events leading to truncated protein products<sup>30</sup>. To date, several AR splice variants have been described, and AR-V7 is the most prevalent and biologically relevant<sup>17</sup>. The first pilot study to demonstrate a potential correlation between AR-V7 and clinical outcomes included 62 patients with mCRPC treated with abiraterone or enzalutamide and assessed the AR-V7 status by a CTC-based RT-PCR assay detecting the AR-V7 messenger RNA<sup>31</sup>. Although preliminary, results demonstrated no PSA declines  $\geq 50\%$  from baseline (PSA<sub>50</sub>) in patients with AR-V7 positive (+) CTCs and also worse PFS and OS compared to AR-V7 negative (-) patients. These results suggested that AR-V7 could potentially serve as a prognostic tool but also possibly a predictive biomarker for response to abiraterone and enzalutamide. Recently, an update of this study with a larger sample size was published, including 202 patients treated with abiraterone or enzalutamide<sup>32</sup>. Importantly, the authors correlated clinical outcomes with 3 different biomarker groups based on CTC detection and AR-V7 detection: CTC(-) vs. CTC(+)/AR-V7(-) vs. CTC(+)/AR-V7(+). Overall, PSA<sub>50</sub> response rates were 75.5% in CTC(-) patients, 52.2% in CTC(+)/AR-V7(-) patients and 13.9% (5 of 36) CTC(+)/AR-V7(+) patients ( $p < 0.001$ ). CTC(+)/AR-V7(+) patients were more likely to have higher Gleason scores ( $\geq 8$ ), prior ARS inhibitor and taxane use, and worse performance status (ECOG  $\geq 1$ ). On multivariable analysis, PFS and OS outcomes were more favorable for CTC(-) patients, intermediate for CTC(+)/AR-V7(-) patients and worse for CTC(+)/AR-V7(+) patients<sup>32</sup>.

In another study conducted in 37 patients with mCRPC treated with taxane-based chemotherapy, no statistical difference in terms of PSA<sub>50</sub>, PFS or OS was detected between AR-V7 (+) vs. AR-V7(-) patients, suggesting that patients may still respond to taxanes regardless of the AR-V7 status<sup>33</sup>. Although a numeric superiority was detected in AR-V7(-) patients in terms of PFS (6.9 vs. 5.1 months,  $P = 0.02$ ) and OS (14.7 vs. 9.2 months,  $P = 0.11$ ), no statistically significant difference was observed. Important to note is that the OS analysis was exploratory and the low number of patients enrolled limits the evaluation of AR-V7 as a prognostic marker in this study. Therefore, it is certainly possible that CTC-based AR-V7 detection using this mRNA assay may be associated with inferior clinical outcomes to chemotherapy if more patients are studied, although the effect is not likely to be as great as in the context of ARS inhibitor therapy.

In a larger study with 161 men with mCRPC treated with ARS inhibitors (abiraterone, enzalutamide or apalutamide) or taxanes (docetaxel, cabazitaxel or paclitaxel), AR-V7 status

was determined using a separate AR-V7 protein immunofluorescent assay performed on a non-EpCAM-based CTC detection platform<sup>34</sup>. This study also demonstrated that patients with AR-V7(+) CTCs treated with ARS inhibitors had worse outcomes by all event measures, including radiographic PFS (2.3 vs. 14.5 months;  $P<0.001$ ), time on therapy (2.1 vs. 6.8 months;  $P<0.001$ ) and OS (4.6 vs. not reached;  $P<0.001$ ). Among patients treated with taxanes, no difference was observed in terms of PFS or time on therapy, but patients with AR-V7 positive CTCs demonstrated inferior OS than AR-V7 negative (8.9 vs. 19.8 months;  $P<0.001$ ), suggesting that AR-V7 status may be prognostic even in taxane-treated patients. Very importantly, in the AR-V7(+) subset, patients treated with taxanes had superior outcomes, including OS, compared to men treated with ARS inhibitors (positive statistical interaction between biomarker status and therapy type), supporting the preliminary evidence of a predictive role of protein-based AR-V7 detection in this setting<sup>34</sup>. In addition, a subsequent study from these same investigators demonstrated that the predictive ability of the biomarker with respect to discriminating the two different types of therapies was greatest when nuclear-localized AR-V7 presence was required to define a biomarker-positive test, rather than any (nuclear or cytoplasmic) presence of AR-V7 protein<sup>35</sup>. For this reason, this assay is now defined as positive only if AR-V7 is nuclear-localized; this definition is now being used in multiple prospective validation studies.

One important limitation of some of the CTC-based AR-V7 detection tests is that a subset of patients may present with AR-low or AR-negative CTCs, often associated with an aggressive phenotype and lack of response to ARS inhibitors, and therefore may be classified as AR-V7 negative. Although this may be a limitation of EPCAM-based CTC assays, CTCs have been identified in AR-low prostate cancer cells using non-EPCAM based tests), including neuroendocrine prostate carcinomas<sup>36</sup>. Also, AR-V7 has been identified in AR-low prostate cancer such as neuroendocrine carcinomas<sup>37</sup>.

In addition to CTC-derived AR-V7 detection, there are other ways to interrogate AR-V7 using liquid biopsies. For example, two recent studies correlated AR-V7 mRNA detection in the whole blood with clinical outcomes in patients with mCRPC treated with abiraterone or enzalutamide<sup>38,39</sup>. The first study, using semi-quantitative mRNA analysis of AR-V7 using analog PCR in peripheral whole blood of 85 men demonstrated worse outcome in patients with high-AR-V7 expression levels, including no PSA<sub>50</sub> responses, shorter PFS and OS<sup>38</sup>. The second study assessed AR-V7 (and PSA) mRNA levels in the whole blood of mCRPC patients using a highly-quantitative digital droplet PCR (ddPCR) methodology, and observed potential prognostic value of both tests on survival of patients treated with ARS inhibitors<sup>39</sup>. A summary of all methods currently used to detect AR-V7 in tumor and liquid biopsies is provided in Table 4.

To add further evidence of the prognostic impact of AR-V7 in patients with mCRPC treated with ARS inhibitors, a meta-analysis on this subject was recently published<sup>40</sup>. In the PFS and OS analysis, a total of eight trials encompassing 490 patients were included and demonstrated that both PFS and OS were better in AR-V7(-) than AR-V7(+) patients. Prospective validation of AR-V7 is now urgently needed.

The first large prospective randomized trial to demonstrate the prognostic significance of AR-V7 status in mCRPC was the ARMOR3-SV trial, which compared galeterone vs. enzalutamide in patients with CTC-based AR-V7(+) disease who had not previously received ARS inhibitors or taxane agents<sup>41</sup>. Galeterone is a selective multi-targeted molecule that acts as a CYP17 inhibitor, AR antagonist and degrader (including AR-V7 degrader), with potential activity in AR-V7(+) patients based on initial preclinical and initial clinical studies<sup>42</sup>. The AR-V7 biomarker test used for eligibility assessment was a derivation of the mRNA-based RT-PCR assay first reported by the Johns Hopkins group<sup>31</sup>. Of the 953 patients screened using this assay, only 8% (73) were AR-V7(+) and 38 patients were randomized to enzalutamide or galeterone. In this trial, AR-V7 detection was associated with worse prognostic features, including higher baseline PSA levels, more bone lesions, higher ECOG PS status, prior anti-androgen and docetaxel use compared to patients with no CTC detected or AR-V7(-) CTCs<sup>41</sup>. Because 7 of the first 12 treated AR-V7(+) patients developed rapid disease progression (early censoring rate of 58%), the data and safety monitoring committee (DSMC) recommended premature termination of the study since it was unlikely to meet its primary endpoint. As a result of this early trial closure, galeterone is no longer being developed as a prostate cancer therapeutic.

Despite all of available data, we must emphasize that further prospective validation studies are important to confirm the role of AR-V7 as a prognostic biomarker in patients with mCRPC. One such study is the PROPHECY trial (NCT02269982). In this trial, 120 mCRPC patients who are beginning next-generation ARS inhibitor therapy will be prospectively sampled at baseline using 3 different AR-V7 liquid biopsies (two are PCR-based mRNA assays, and one is a immunofluorescence-based protein assay), and then patients will be monitored using a prospectively-defined follow-up schema until clinical or radiographic progression of their disease. At the time of progression, patients will be re-sampled using the same 3 liquid biopsy platforms, and will then be offered a taxane chemotherapy. At the time of progression on the taxane agent, a third set of liquid biopsy samples for AR-V7 analysis will be collected. At the time of writing, this study has completed enrolment of all 120 men, and preliminary data on the primary endpoint of PFS are eagerly awaited. One of the unique advantages of this study is its ability to assess the prognostic value of each of the 3 AR-V7 assays, and to indirectly compare the analytical and clinical characteristics of each test against the others. A preliminary communication of the top-line results from the PROPHECY trial is expected by Q2 2018.

#### 4. POTENTIAL TOOLS FOR TREATMENT SELECTION

An important unmet need for patients with metastatic CRPC is the identification of biomarkers that can be predictive for response to available therapies, to guide treatment selection at each specific time point in the course of disease, with the ultimate goal of maximizing outcomes and limiting toxicity. Although there are no currently available validated predictive biomarkers in this setting, many are under development and hopefully in the near future will help us to select the right therapy for the right patient at the right time.

AR-V7 is one of these candidates as a putative predictor of benefit to the newer ARS inhibitors, as demonstrated in several single-center trials and a recent meta-analysis, as

discussed above<sup>31,32,34,38,40</sup>. These studies demonstrate in aggregate that the benefit of ARS inhibitors is limited to AR-V7(-) patients and very few patients with AR-V7(+) disease demonstrate benefit with abiraterone or enzalutamide. On the other hand, it appears that AR-V7 status does not interfere as dramatically with responsiveness to taxane-based chemotherapy<sup>33,34</sup>. Therefore, AR-V7 status may represent a marker for therapy selection in mCRPC if larger studies confirm its predictive role in this scenario. Recently, two studies presented data suggesting that some patients with AR-V7(+) may achieve PSA declines with enzalutamide or abiraterone, although a transient PSA reduction does not always equate with a meaningful clinical benefit. The disappointing results of the ARMOR3-SV study comparing galeterone vs. enzalutamide in AR-V7(+) patients were recently reported as discussed above, also showing some PSA<sub>50</sub> responses to enzalutamide<sup>41</sup>. Another recent study described that 6 of 21 AR-V7(+) patients demonstrated PSA reductions under treatment with abiraterone or enzalutamide, but again this does not necessarily equate with clinical benefit<sup>43</sup>. Despite several discussion points on the interpretation of these data<sup>44,45</sup>, specially related to benefit demonstrated only by PSA decline and no other outcome endpoints, the bottom of line is that large prospective validation studies are critical to define the real impact of AR-V7 status on treatment selection between ARS inhibitors and taxane chemotherapy. Interestingly however, a recent study from the Johns Hopkins group evaluating the real-world clinical utility of AR-V7 testing using a commercial CLIA-certified assay suggested that physicians used the AR-V7 results more often than not to inform their clinical decisions, and that the physicians who used the AR-V7 test results to direct the next line of systemic therapy observed greater rates of PSA<sub>50</sub> declines with the subsequent systemic therapy compared to physicians who did not use the results to change treatment<sup>46</sup>. These provocative but preliminary findings suggest that AR-V7 testing is already beginning to demonstrate broader clinical utility outside of specialized academic centers.

Other potential predictive biomarkers under investigation for mCRPC therapy selection are: (1) AR mutations detected by cell-free DNA; (2) DNA damage repair gene alterations as predictive for response to PARP inhibitors; (3) gene alterations related to aggressive prostate cancer phenotype and response to platinum agents; and (4) microsatellite instability (MSI) as a predictor for benefit to pembrolizumab, an anti-PD inhibitor. In the near future, these and possibly others markers may help us to select the best treatments and avoid futile therapies for patients with metastatic castration-resistant prostate cancer. A list of these and other putative predictive markers under investigation is included in Table 5.

## 5. CONCLUSIONS

There is a growing body of evidence supporting blood-based AR-V7 status as a prognostic biomarker in metastatic castration-resistant prostate cancer. Patients with AR-V7(+) disease tend to present with more aggressive features, evidenced by higher PSA levels, higher disease burden and worse performance status. The predictive role of the AR-V7 biomarker to aid in treatment selection between ARS inhibitors and taxane agents requires further validation, but preliminary evidence to date suggests limited efficacy of abiraterone or enzalutamide in AR-V7(+) patients while taxanes appear to retain some level of sensitivity despite AR-V7 detection. While definitive prospective trials (such as PROPHECY) are

awaited, existing data point to some evidence of clinical utility of the AR-V7 biomarker in real-world oncology practices. The era of precision medicine for prostate cancer is very near!

## 6. Expert commentary

Recent advances in the understanding of prostate cancer biology and mechanisms of progression to androgen deprivation therapy have led to the development of several biomarkers that can be used to assess prognostic and also potentially predictive biomarkers which could serve as treatment-selection tools. To date, there are many prognostic biomarkers identified in prospective studies, including lactate dehydrogenase (LDH) levels, ECOG performance status, PSA level, sites of disease, hemoglobin, baseline CTC count (  $\geq 5$ ) and decline (to  $< 5$ ) with therapy, among others. Recently, AR-V7 status has been correlated with clinical outcomes and patients with AR-V7 detected in liquid biopsies tend to present with more aggressive disease and shorter progression-free and overall survival.

As discussed herein, an important unmet need for patients with metastatic castration-resistant prostate cancer (mCRPC) is the identification of predictive biomarkers that could be used by oncologists to recommend therapies with higher likelihood of response and to avoid ineffective drugs, maximizing clinical benefits to patients while minimizing toxicities. To date, there is a growing body of evidence supporting AR-V7 status as a potential tool in this setting, since studies have demonstrated a lack of significant benefit with novel AR signaling inhibitors in AR-V7 positive patients, but partial sensitivity to taxane-based chemotherapy that may potentially occur in the context of AR-V7 conversions from positive to negative. Although these findings still require validation in larger studies, recent data indicate that AR-V7 status may have an important role to guide treatment decisions in clinical practice. Another promising biomarker under investigation that could be integrated together with the assessment of the AR-V7 status is the presence of specific AR mutations, which also can help to guide the best choice of systemic therapies. For example, a patient with mCRPC receiving abiraterone with a rising PSA who is found using a liquid biopsy to harbor the AR T878A mutation (associated with abiraterone resistance) could be switched to enzalutamide early before developing clinical progression on abiraterone. If that same patient presented with AR-V7 positive CTCs as well, it is unlikely that he would respond to enzalutamide and would probably be best served with alternative agents including docetaxel, cabazitaxel, or even radium-223.

There are many other promising biomarkers under investigation beyond AR-V7 and AR mutations, including AR-independent pathways such as the presence of DNA damage response (DDR) gene alterations, which may predict benefit to PARP inhibitors and platinum chemotherapies, the presence of microsatellite instability which might predict benefit to immunotherapy (anti-PD1 agents, *i.e.* pembrolizumab) and other gene alterations such as Rb loss and TP53 mutations, which have been associated with an aggressive phenotype and poor response to AR-directed therapies. We expect that the incorporation of these and other biomarkers in the sequential assessment of patients with advanced prostate cancer - *i.e.* precision medicine - will lead to a more rational use of available and future therapies, with significant improvements in outcomes for our patients.



## 7. Five-year view

We believe that in the near future therapy selection for patients with metastatic castration-resistant prostate cancer will be guided by predictive biomarkers identified in liquid biopsies and/or tumor tissue. It is likely that many of the potential biomarkers under investigation will be integrated into routine clinical practice, including assessment of AR-V7 status, AR mutations, CTC heterogeneity, presence of DNA repair gene alterations, microsatellite instability, and potentially others. Moreover, a better understanding of the disease biology in an individual patient at a particular moment in time will help to integrate other strategies, such as potentially local therapy to the prostate gland and also rational drug combination strategies. This will likely result in increased efficacy and value of the available therapies, ultimately leading to improvements in important endpoints such as survival and quality of life.

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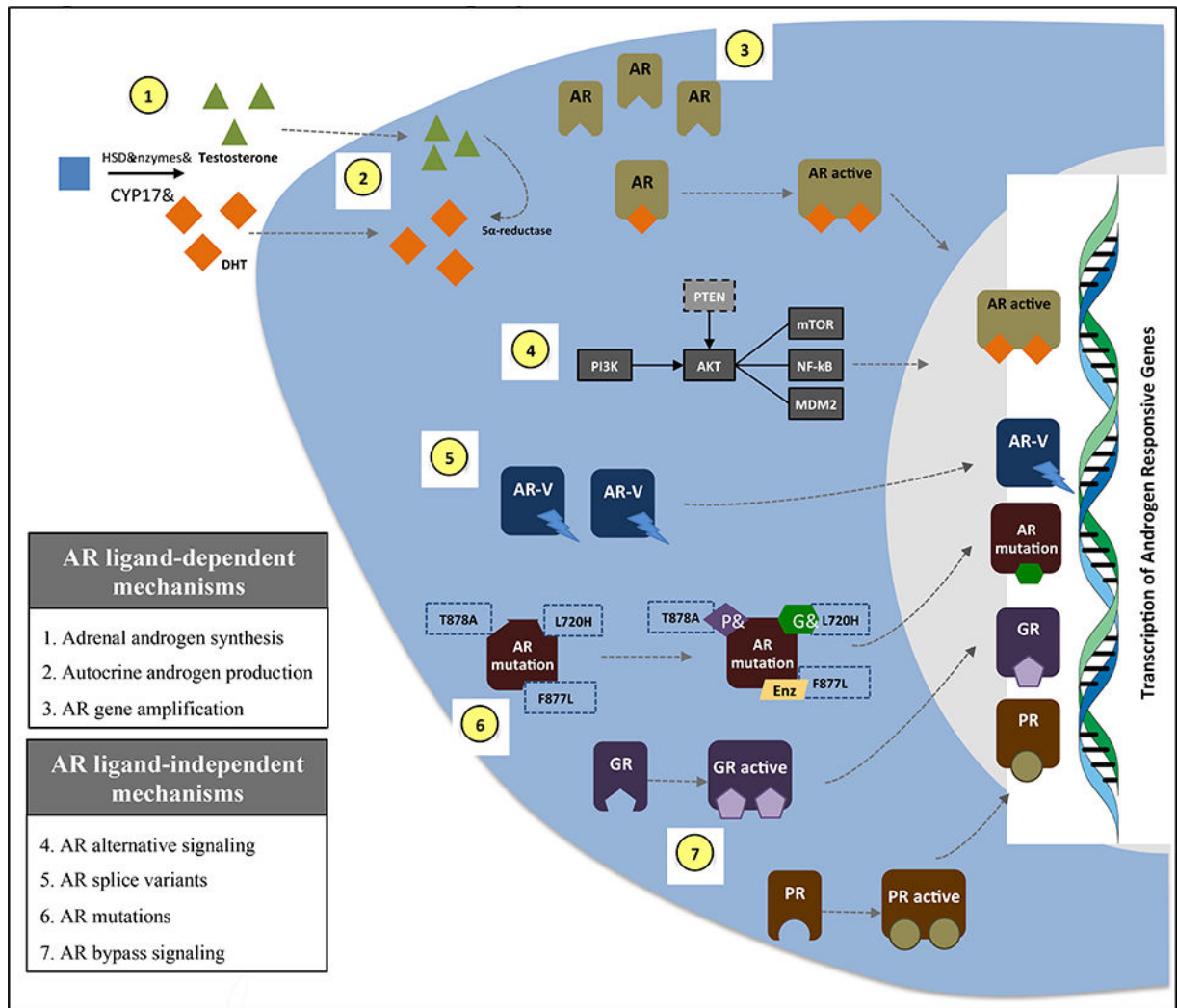
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### Key issues

- Currently, there are several life-prolonging therapies available for patients with mCRPC, but treatment selection is largely based on clinical factors and no validated predictive biomarkers are yet available to guide therapeutic choices.
- The mechanisms of escape to novel androgen-receptor signaling (ARS) inhibitors are now better understood and include ligand-dependent and -independent mechanisms.
- mCRPC remains a highly heterogeneous disease and many prognostic biomarkers have been identified to help estimate survival in this setting, such as lactate dehydrogenase (LDH), hemoglobin, ECOG performance status, PSA, albumin, and others.
- Recently, CTC count and AR-V7 status have been described as prognostic markers. The presence of a baseline CTC count  $\geq 5$  (and post-treatment CTC conversion to  $<5$ ) and/or AR-V7 detection are associated with overall worse survival.
- There have been significant advances to identify biomarkers for therapy selection, such as AR-V7 status, AR mutations, CTC heterogeneity, the presence of DNA damage repair gene alterations. Prospective validation of these biomarkers is still pending prior to incorporation in routine clinical use.
- AR-V7 is a constitutionally active form of the AR and has been implicated in primary and acquired resistance to ARS inhibitors such as abiraterone and enzalutamide, but is compatible with sensitivity to taxane chemotherapies.
- Patients with AR-V7-positive CTCs may still respond to taxane-based chemotherapy, including some cases that are associated with AR-V7 conversions from positive to negative. Therefore, if these findings are confirmed in validation studies, AR-V7 status may serve as a treatment-selection tool for patients with mCRPC.



**Figure 1. Androgen receptor signaling axis and potential androgen-dependent and independent mechanisms of disease progression.**

Figure 1 highlights the androgen receptor (AR)-signaling axis, with conversion of testosterone to dihydrotestosterone (DHT) by the 5 $\alpha$ -reductase enzyme, and subsequent AR activation, dimerization, nuclear translocation and activation of transcriptional activation of target genes. The figure summarizes potential mechanisms of resistance to AR-signaling inhibitors by using a schematic representation of a prostate cancer cell. *Not shown* are multiple additional androgen/AR-independent mechanisms of escape including activated Wnt pathway signaling, loss of the RB1 and/or TP53 genes, overexpression of DNA repair pathways proteins including PARP1 and DNA-PK, epigenetic dysregulation (*e.g.* via EZH2 overexpression), and neuroendocrine/small cell transformation.

**Abbreviations:** CYP17: cytochrome P450 17 $\alpha$ -hydroxylase; PI3K: phosphoinositide 3-kinase; AKT: protein kinase B; PTEN: phosphatase and tensin homolog; mTOR: mammalian target of rapamycin; NF- $\kappa$ B: nuclear factor kappa B; MDM2: mouse double minute 2 homolog; AR-V: AR splice variant; P: progesterone; PR: progesterone receptor; G: glucocorticoid; GR: glucocorticoid receptor.



**Table 1.**

Current systemic treatment options for metastatic prostate cancer.

<b>CASTRATION-SENSITIVE PROSTATE CANCER</b>	
<b>ADT</b> •LHRH agonists vs. antagonists •Continuous vs. intermittent	<ul style="list-style-type: none"> <li>•Mainstay of therapy for mCSPC.</li> <li>•No clear superiority of LHRH agonists or antagonists.</li> <li>•Intermittent ADT not proven to be non-inferior to continuous<sup>47</sup>.</li> <li>•Should be used continuously in the setting of mCRPC.</li> </ul>
<b>ADT plus Docetaxel</b>	<ul style="list-style-type: none"> <li>• Two RCTs and meta-analysis demonstrating improved outcomes for de novo mCSPC, including OS<sup>12,13,48</sup>.</li> </ul>
<b>ADT plus Abiraterone and prednisone</b>	<ul style="list-style-type: none"> <li>• Two RCTs demonstrating significant improvements in outcomes for de novo mCSPC, including OS<sup>14,15</sup>.</li> </ul>
<b>CASTRATION-RESISTANT PROSTATE CANCER</b>	
<b>AR-signaling inhibitors</b> •Abiraterone / prednisone •Enzalutamide	<ul style="list-style-type: none"> <li>•Both with RCTs demonstrating significant improvements in OS and QoL in pre- and post-chemo settings<sup>4-7</sup>.</li> <li>•Sequential use not proven to improve clinically meaningful outcomes<sup>49</sup>.</li> </ul>
<b>Taxane-based chemotherapy</b> •Docetaxel •Cabazitaxel	<ul style="list-style-type: none"> <li>•RTCs demonstrating improvements in clinically meaningful outcomes<sup>8,9</sup>.</li> <li>•Cabazitaxel approved for patients previously treated with docetaxel<sup>9</sup>.</li> <li>•Both taxanes retain activity in patients with AR-V7<sup>33,34</sup>.</li> </ul>
<b>Radiopharmaceuticals</b> • Radium-223	<ul style="list-style-type: none"> <li>•1<sup>st</sup> radiopharmaceutical to demonstrate OS benefit<sup>10</sup>.</li> <li>•Can be used in pre- and post-chemo settings, and in combination with osteoclast-inhibitory agents.</li> </ul>
<b>Immunotherapy</b> • Sipuleucel-T	<ul style="list-style-type: none"> <li>• Despite OS improvement demonstrated in one RCT, no benefit in terms of PSA decline, objective response or PFS<sup>11</sup>.</li> </ul>

Abbreviations: ADT: androgen deprivation therapy; aPC: advanced prostate cancer; CRPC: castration-resistant prostate cancer; RCT: randomized clinical trials; OS: overall survival; QoL: quality of life; PFS: progression-free survival; chemo: chemotherapy; PSA: prostatic specific antigen.

**Table 2.**

Biomarker category definitions<sup>19</sup> and examples in advanced prostate cancer.

<b>Biomarker Category</b>	<b>Definition</b>	<b>Examples</b>
<b>Diagnostic</b>	Used to confirm presence of disease or identify disease subtype.	Pathology (prostate biopsy).
<b>Monitoring</b>	Used serially to assess status or extent of a disease.	PSA, bone scan, CTC count.
<b>Response</b>	Used to show a biologic response to an agent or product.	PSA, Bone scan, CTC conversion.
<b>Prognostic</b>	Identifies at baseline the likelihood of a clinical event (e.g., progression, death), irrespective of treatment.	Performance status, LDH, visceral disease, presence of pain, etc.
<b>Predictive</b>	Identifies individuals at baseline who are more likely to experience a favorable or unfavorable effect from one agent compared to another.	None validated so far. Potential candidates are AR-V7, CTC heterogeneity, and DDR gene alterations.
<b>Safety</b>	Indicates the likelihood or extent of toxicity as an adverse event of an agent.	Performance status, hepatic function for some therapies, renal function for other therapies.
<b>Susceptibility</b>	Indicates the potential risk for developing a medical condition.	Family history, germline BRCA2 mutation, African American race.

Abbreviations: PSA: prostatic specific antigen; LDH: lactate dehydrogenase; AR-V7: androgen receptor splice variant 7; MSI: microsatellite instability; DDR: DNA damage repair.

**Table 3.**

Key prognostic biomarkers in metastatic prostate cancer.

Laboratory biomarkers <sup>22</sup>	Clinical biomarkers <sup>22</sup>
- PSA (baseline and kinetics) -LDH - Hemoglobin level - Alkaline phosphatase - Serum albumin - CTC count 5 vs. <5 at baseline and CTC count conversion (<5) after 12 weeks of therapy <sup>23-26</sup>	- Performance status - Presence of pain / use of opioids
	<b>Imaging biomarkers</b>
	- Sites of metastasis <sup>50</sup> - Extent of disease <sup>12,51</sup>
	<b>Pathologic / Molecular biomarkers</b>
	- Aggressive histologic variants (e.g. small cell) - Rb1, TP53 and/or PTEN alterations <sup>52</sup> . AR-V7+ detection <sup>32,40</sup> DDR gene alterations

Abbreviations: PSA: prostatic specific antigen; LDH: lactate dehydrogenase; CTC: circulating tumor cell; AR-V7 androgen receptor splice variant 7.

**Table 4.**

Currently used methods to detect AR-V7 in tumor and liquid biopsies

<b>Tissue-based AR-V7 detection methods</b>
<ul style="list-style-type: none"> <li>- Western blot for protein detection<sup>53</sup></li> <li>- IHC for protein detection<sup>54</sup></li> <li>- RNA in situ hybridization (RISH) for mRNA<sup>55-57</sup></li> <li>- RT PCR for mRNA detection<sup>58</sup></li> <li>- RNA sequencing for mRNA detection<sup>59</sup></li> </ul>
<b>Blood-based AR-V7 detection methods</b>
<ul style="list-style-type: none"> <li>- CTC based RT PCR for mRNA detection<sup>31</sup></li> <li>- CTC based ddPCR for mRNA detection<sup>60</sup></li> <li>- CTC based RNA-seq<sup>61</sup></li> <li>- CTC based protein (immunofluorescence] detection<sup>34</sup></li> <li>- Whole blood analog RT PCR for mRNA detection<sup>38</sup></li> <li>- Whole blood ddPCR for mRNA detection<sup>39</sup></li> <li>- Cell free RNA detection by RT-PCR<sup>62</sup></li> <li>- Exosome derived PCR detection of mRNA<sup>63</sup></li> </ul>

Abbreviations: AR-V7: androgen receptor splice variant 7; IHC: immunohistochemistry; RT PCR: reverse transcription polymerase chain reaction; mRNA: messenger ribonucleic acid.

**Table 5.**

Potential predictive biomarkers under investigation for mCRPC therapy-selection.

Potential Predictive Biomarker	Therapy Selection
AR-V7 <sup>31,32,34,64</sup>	Not detected: ARS inhibitors Detected: taxane chemotherapy
Activating AR-LBD mutations and/or <i>AR</i> gene amplification <sup>65,66</sup>	Not detected: ARS inhibitors Detected: taxane chemotherapy
CTC heterogeneity <sup>67</sup> (Shannon index)	Low (Shannon <1.5): TARS inhibitors High (Shannon ≥ 1.5): taxane chemotherapy
DDR gene alterations	PARP inhibitors <sup>68</sup> or platinum agents <sup>69</sup>
Microsatellite instability (MSI-high)	Immunotherapy with anti-PD-1 (pembrolizumab) <sup>70</sup>

Abbreviations: ARS: androgen receptor signaling; AR-V7: androgen receptor splice variant 7; AR-LBD: androgen receptor ligand-binding domain; CTC: circulating tumor cell; DDR: DNA damage repair; PARP: poly ADP (*adenosine diphosphate*)-ribose polymerase; PD-1 programmed death.