



Review

Metastasis in renal cell carcinoma: Biology and implications for therapy



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Abstract Although multiple advances have been made in systemic therapy for renal cell carcinoma (RCC), metastatic RCC remains incurable. In the current review, we focus on the underlying biology of RCC and plausible mechanisms of metastasis. We further outline evolving strategies to combat metastasis through adjuvant therapy. Finally, we discuss clinical patterns of metastasis in RCC and how distinct systemic therapy approaches may be considered based on the anatomic location of metastasis.

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1. Introduction

Approximately one-third of patients with renal cell carcinoma (RCC) present with metastatic disease, and amongst those patients with localized disease, a substantial proportion will recur [1]. For patients with metastatic renal

cell carcinoma (mRCC), the landscape of therapy has evolved dramatically over the past decade. Prior to 2005, immunotherapy represented the mainstay of therapy with agents such as interleukin-2 (IL-2) and interferon- α (IFN- α) [2,3]. Estimates of overall survival (OS) in that era coalesced around 1 year. Since 2005, multiple targeted therapies have been approved, primarily directed at vascular endothelial growth factor (VEGF) or its cognate receptor (VEGF receptor, or VEGFR), or the mammalian target of rapamycin (mTOR) [4]. In this generation of therapies, inhibitors of VEGF include axitinib, bevacizumab, pazopanib, sorafenib and sunitinib, while inhibitors of mTOR include

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everolimus and temsirolimus [5–10]. These agents have collectively improved median survival estimates to approximately 2.5–3 years [11]. Over the past year, 3 additional FDA approvals have been granted for mRCC for a VEGFR/MET/AXL inhibitor (cabozantinib), a programmed death-1 (PD-1) inhibitor (nivolumab) and a multikinase inhibitor (lenvatinib, approved with everolimus) [12–14]. It remains to be seen how these agents will alter OS estimates for mRCC, although it will surely move the bar in a positive direction.

Despite these critical advances, the reality is that the vast majority of patients with mRCC have incurable disease [15]. A goal of treatment is to maximize the yield of existing systemic therapies through personalized approaches. In the current review, we explore how clinical and biological properties of metastases may potentially alter paradigms for systemic therapy.

2. Biology of RCC

2.1. Differing biology by histology

It is critical to acknowledge that mRCC is comprised of multiple distinct histologies, each with unique biologic underpinnings. The most common histology is clear cell, comprising 75%–80% of cases. Approximately 70% of patients with clear cell RCC bear alterations in the Von-Hippel Lindau (*VHL*) gene [16]. Wild type *VHL* protein functions as an ubiquitin ligase, participating in degradation of hypoxia inducible factor (HIF). In patients bearing *VHL* alteration, the resulting high levels of HIF result in upregulation of VEGF. VEGF activates VEGFR, triggering the phosphoinositol-3-kinase (PI3K)-Akt signaling cascade. Downstream, mTOR is activated and leads to transcription of a variety of tumor-promoting factors, resulting in increased cellular migration and angiogenesis. Although VEGFR has typically been implicated as the key driver of tumor progression in RCC, there is emerging evidence that other transmembrane receptors may potentially drive metastasis, including MET and AXL [17].

Non-clear cell RCC histologies comprise roughly 20%–25% of patients overall. The most prevalent of these is papillary RCC, which represents 10%–15% of patients. Papillary RCC is frequently subdivided into type I and type II disease. Type I disease is characterized by alterations in the MET proto-oncogene, while type II is characterized by a variety of alterations. Recent data from The Cancer Genome Atlas (TCGA) investigators highlighted alterations in *SETD2*, *CKDN2A* and *TFE3* fusions as frequent events in type II papillary RCC [18]. Chromophobe type disease comprises approximately 5% of all RCC cases. TCGA data pertaining to chromophobe RCC suggest frequent changes in the *TERT* promoter region, and mitochondrial DNA analyses suggest changes in mitochondrial function [19]. Beyond papillary and chromophobe RCC, other histologies of RCC represent <1% of all cases. Despite their rarity, there are efforts to characterize the genomic changes occurring in these entities. For instance, our group has identified frequent alterations in *NF2* in patients with collecting duct RCC, an exquisitely rare diagnosis with a dismal prognosis [20].

Admixed with any histological subset of RCC may be sarcomatoid elements. Sarcomatoid RCC is thought to

coexist with other histologies in about 25% of cases [21]. Sarcomatoid disease tends to be particularly aggressive, although (as discussed subsequently) the current treatment paradigm is not distinct from clear cell disease. Our group has identified frequent alterations in the aurora kinase pathway, and *NF2* alterations have also been detected in this disease [22,23].

2.2. Tumor heterogeneity

Although histology is frequently used to offer prognostic data to patients, it is critical to acknowledge that the biology of tumors may differ across sites of metastasis. One of the first detailed studies to identify this intratumoral heterogeneity was from Gerlinger and colleagues [24]. In an effort that included just 4 patients with mRCC, separate sites of metastasis were evaluated. Alterations in the mTOR pathway were variable across sites of metastasis, as were alterations in *SETD2*, *PTEN* and *KDM5C*. Subsequent sections will highlight potential therapeutic strategies for these alterations. With the evolution of novel immunotherapeutic strategies, there has also been substantial interest in characterizing PD-L1 expression in metastatic sites. A recent study from the Dana Farber Cancer Institute compared tissues derived from 53 primary RCC specimens and 73 corresponding metastases [25]. PD-L1 expression appeared to be consistent, although PD-L1 expression was noted to be heterogeneous within lesions.

2.2.1. Biological mediators of metastasis

Little is agreed upon regarding the biological mechanisms that drive RCC metastasis. On a macromolecular level, Grange et al. [26] have proposed that tumor-derived microvesicles (which essentially break off from the primary site) may disperse tumors through hematogenous routes. These microvesicles appear to bear CD105-positive cells, which carry a cancer stem cell phenotype, and microRNAs which stimulate angiogenesis. The immune milieu may also play a critical role in the evolution of metastases. In preclinical models, neutrophilic infiltration in the lungs (accompanied by secretion of neutrophil chemokines) was accompanied by suppression of pulmonary metastases of RCC [27]. In contrast, loss of neutrophil chemokines in the lung was accompanied by an increase in pulmonary metastases. Other immune cells with negative effects on antitumor immunity (e.g., myeloid derived suppressor cells, or MDSCs) have been shown to have a pro-angiogenic effect and cause propagation of RCC in preclinical models [28].

Beyond these macromolecular events, several molecular mediators of RCC metastasis have been identified. In the setting of clear cell RCC bearing *VHL* alteration, it has been proposed that CUB-domain-containing protein (CDCP1) may drive metastasis [29]. CDCP1 is regulated through HIF dependent pathways and drives activation of protein kinase C- δ (PKC δ), which in turn increases cellular migration. Expression of MUC1, a membrane-bound glycoprotein, is also HIF-dependent, and knockdown of MUC1 has been shown to markedly decrease cellular invasion and migration in *in vitro* RCC models [30]. Various chemokine receptors, including CXCR4, also appear to be upregulated in the

context of clear cell RCC. An increased expression of CXCR4 and its ligand CXCL12 in the setting of *VHL* alteration has been associated with increased metastatic spread in pre-clinical studies [31].

3. Prevention of tumor metastasis: role of adjuvant therapy

3.1. Localized disease

At present, there is no defined role for adjuvant therapy for localized RCC following partial or radical nephrectomy. The current standard of care involves serial imaging with computerized tomography, typically up to a span of 5 years. However, multiple trials have evaluated the strategy of using approved agents in the metastatic setting as adjuvant therapy [32]. Both adjuvant VEGF-tyrosine kinase inhibitors (VEGF-TKIs) and mTOR inhibitors have been evaluated, including sunitinib, sorafenib, pazopanib, axitinib and everolimus.

The first phase III study to report out exploring adjuvant therapy is the phase III ASSURE trial [33]. This study randomized 1943 patients to receive one of two VEGF-TKIs (sunitinib or sorafenib) or placebo. The study had to be modified on the basis of early toxicity events; in a modification to the original design, patients were started on a lower dose of sunitinib and sorafenib. Doses were escalated if no toxicities were incurred after a 2-month period. The primary endpoint of the study was disease-free survival, and the trial identified no significant difference across the three cohorts (5.8 years with sunitinib, 6.1 years with sorafenib and 6.6 years with placebo). Aside from failing to meet the primary endpoint of the study, there were also concerns regarding treatment related toxicity. Toxicities encountered were typical for VEGF-directed therapies, with the most common adverse events being hypertension, hand-foot syndrome, rash and fatigue. A total of five deaths occurred within the first month of protocol-based therapy. Four of these patients had received sunitinib, with deaths attributable to neurologic dysfunction, gastrointestinal perforation, pulmonary embolus and disease progression, respectively. The remaining patient received sorafenib and developed infectious colitis during therapy. In the adjuvant setting, even if a modest benefit with adjuvant therapy was observed, it would need to be counterbalanced against the toxicity profile and the potential risk of treatment-related mortality.

The remaining trials of adjuvant targeted therapy in RCC have yet to report out. Each of these studies includes a slightly different patient population and treatment strategy, and thus, it is possible that the results may differ from the aforementioned ASSURE trial (Table 1). For instance, the phase III ATLAS study compares a total of 3 years of axitinib therapy to placebo, and the phase III SORCE study similarly examines 3 years of adjuvant sorafenib. The phase III EVEREST trial uniquely explores the mTOR inhibitor everolimus – it is possible that the increased tolerability of mTOR inhibition may make enhance adherence and dose intensity. Although ASSURE is the first study to report final results, data from the phase III S-TRAC trial are anticipated shortly. A recent press release suggested that this trial (a

comparison of 1 year of adjuvant therapy with sunitinib to placebo) met its primary endpoint of improved disease-free survival [34]. The study differed from ASSURE in that it included a higher risk population. Specifically, while ASSURE allowed enrollment of patients with pT1b and grade 3–4 disease, or pT2-4 disease and any grade, S-TRAC enrollment was limited to patients with pT3-4 disease with any grade. The study included just over 670 patients, a sample size considerably smaller than ASSURE.

It is curious that S-TRAC would yield a positive result. A subset analysis of the ASSURE study limited to pT3-4 patients still failed to show a benefit in disease-free survival, despite including a sample size comparable to the overall S-TRAC study population. Subtle differences such as radiographic interpretation may potentially account for the discordant results – S-TRAC employed central radiographic review, while local review was performed in the ASSURE trial. In any case, the implementation of results from S-TRAC will likely be contingent on a number of factors as yet unknown, including the magnitude of benefit with adjuvant sunitinib in this study and the degree of toxicity incurred.

Beyond VEGF- and mTOR-inhibitors, the favorable toxicity profile of PD-1 inhibitors makes these agents attractive for use in the adjuvant setting. Furthermore, as previously noted, studies have shown relatively balanced expression of PD-L1 in both primary tumor and metastases. The Eastern Cooperative Oncology Group (ECOG) has unveiled a plan for assessing both neoadjuvant and adjuvant therapy with nivolumab in a phase III clinical trial. Other agents such as atezolizumab, a PD-L1 inhibitor, may soon be explored in this setting.

3.1.1. Metastatic disease

Metastasectomy is the standard of care for patients who have limited sites of metastasis from RCC. There are limited adjuvant studies that incorporate patients who have had metastasectomy for stage IV RCC – ostensibly, these patients carry the highest potential risk of recurrence, with some retrospective series suggesting a median time to recurrence as low as 16 months [35]. The standard of care in this setting is vigilant observation – although no guidelines exist, monitoring every 3 months with computerized tomography represents a reasonable practice. ECOG 2810 is a prospective, phase III study randomizing patients with fully resected clear cell mRCC to either pazopanib or placebo for 1 year. The study will accrue a total of 126 patients. Previous trials of adjuvant therapy, such as ASSURE, have been criticized for inclusion of low-risk patients, which thereby increases the necessary effect size for any intervention. The high risk of recurrence in the ECOG 2810 study population may poise the trial well for a positive outcome.

4. Patterns of metastases in RCC

One of the most widely cited studies of RCC metastasis distribution is derived from the National Inpatient Sample [36]. In this study, 11,157 patients with mRCC were identified from 1998 to 2007. The most common sites of metastases were lung (45%), following by bone (30%) and lymph node (22%). Liver metastases were noted in 20% of

Table 1 Trials of adjuvant targeted therapy in RCC.

| Trial (sponsor) | Randomization | Treatment duration (year) | <i>n</i> | Limited to clear cell | Inclusion criteria | Details |
|--------------------------|-------------------------------------|---------------------------|----------|-----------------------|--|---|
| <i>Reported trials</i> | | | | | | |
| ASSURE (ECOG) | Sunitinib vs. sorafenib vs. placebo | 1 | 1943 | No | pT1b (G3-4) or pT2-4 or N ⁺ | <ul style="list-style-type: none"> • Study showed no significant difference in DFS across arms |
| S-TRAC (Pfizer) | Sunitinib vs. placebo | 1 | 670 | Yes | pT3-4 or N ⁺ | <ul style="list-style-type: none"> • Press release indicates study met primary endpoint • Further study results to be presented at upcoming meeting |
| <i>Unreported trials</i> | | | | | | |
| ATLAS (Pfizer) | Axitinib vs. placebo | 3 | 592 | Yes | pT2-4 or N ⁺ | <ul style="list-style-type: none"> • Study employs a longer duration of axitinib therapy • Small sample size relative to other adjuvant trials |
| EVEREST (SWOG) | Everolimus vs. placebo | 1 | 1218 | No | pT1b (G3-4) or pT2-4 or N ⁺ | <ul style="list-style-type: none"> • Study nearing completion of accrual • Only adjuvant trial to assess an mTOR inhibitor |
| PROTECT (GSK) | Pazopanib vs. placebo | 1 | 1500 | Yes | pT1b (G3-4) or pT2-4 or N ⁺ | <ul style="list-style-type: none"> • Projected accrual per arm (<i>n</i> = 750) makes this the most sizeable exploration of a VEGF-TKI in a single study |
| SORCE (MRC) | Sorafenib vs. placebo | 3 | 1420 | No | Leibovich score 3-11 | <ul style="list-style-type: none"> • Only study to utilize Leibovich scoring criteria • Leibovich scoring utilizes pathologic elements such as tumor necrosis, which are subject to variability amongst reviewers |

ECOG, Eastern Cooperative Oncology Group; DFS, disease-free survival; mTOR, mammalian target of rapamycin; GSK, GlaxoSmithKline; VEGF-TKI, vascular endothelial growth factor tyrosine kinase inhibitor; MRC, Medical Research Council; RCC, renal cell carcinoma.

patients and adrenal metastases were noted in 9% of patients. Brain metastases occurred in approximately 9% of patients (Fig. 1).

Several series have pointed towards distinct outcomes depending on patterns of metastasis. In an assessment of 2027 mRCC patients from the International mRCC Database Consortium (IMDC) experience, liver metastases were noted to occur in a larger proportion of patients with poor-risk by IMDC criteria [37]. Furthermore, the hazard ratio for death (adjusted for IMDC risk factors) was 1.4 for patients with bone metastases (95%CI 1.22–1.62) and 1.42 for patients with liver metastases (95%CI 1.17–1.73). IMDC data have also been used to assess the outcome of patients with brain metastases [38]. Approximately 106 patients were identified with brain metastases, of which only 12% fell into the favorable IMDC risk group. Most patients (90%) had cerebral metastases, while a smaller subset (17%) had cerebellar metastases. As one might anticipate, a higher number of brain metastases were associated with poorer outcome.

There are several sites of metastases that are associated with more favorable prognoses. Multiple anecdotal reports exist documenting regression of lung metastases following either brief local or systemic therapy for mRCC, and broadly, patients with lung-only metastases are thought to carry a favorable prognosis [39–41]. Interestingly, pancreatic metastases are also thought to confer a favorable prognosis. In a retrospective series from MD Anderson, 228 patients with pancreatic metastases were characterized. Median OS was 39 months in patients with pancreatic metastases as compared to 26 months in patients without ($p < 0.010$).

5. Treatment by anatomic site

5.1. Bone metastases

While a comprehensive discussion of RCC treatment options is beyond the scope of this review, we discuss herein unique

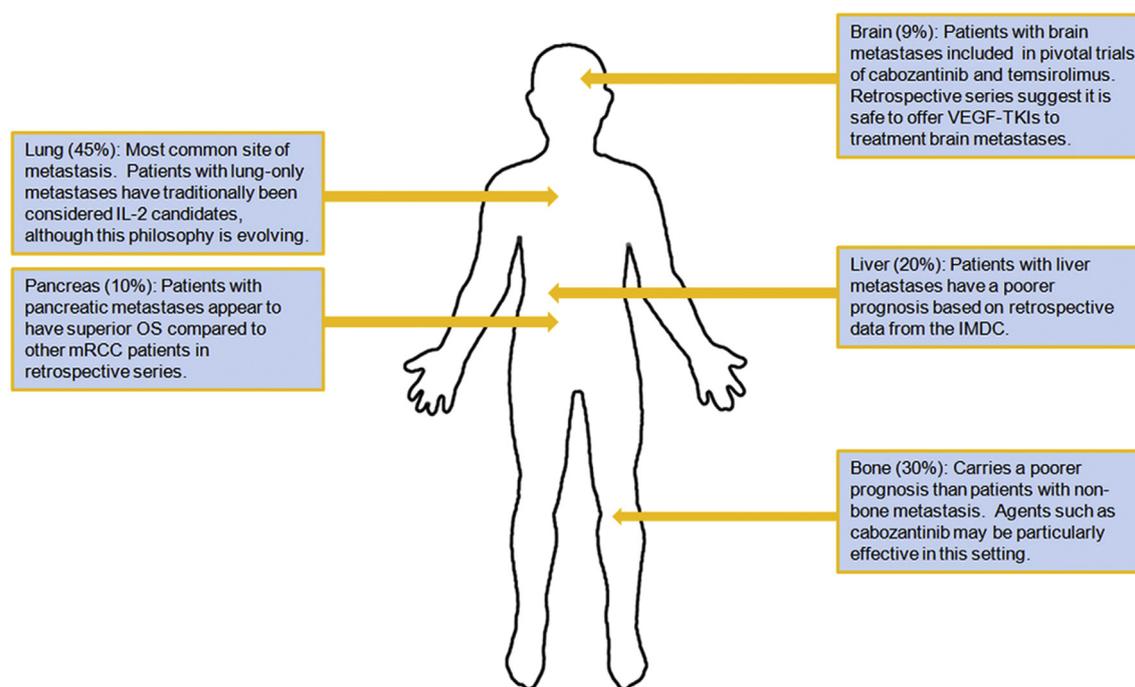


Figure 1 Unique considerations for sites of metastasis in mRCC. IL-2, interleukin-2; IMDC, International mRCC Database Consortium; mRCC, metastatic renal cell carcinoma; OS, overall survival; VEGF-TKIs, VEGF-tyrosine kinase inhibitors.

systemic therapy considerations that pertain to specific anatomic sites. In particular, bone and brain metastases represent significant therapeutic dilemmas. For patients with bone metastases, recent data have alluded to the potential utility of cabozantinib. Cabozantinib, a multikinase inhibitor with activity against VEGFR, MET and AXL, has previously been demonstrated to have impressive activity in bone metastases derived from prostate cancer. In this setting, disappearance of lesions on bone scan has been observed [42]. In mRCC, cabozantinib was compared to everolimus in a phase III clinical trial, demonstrating an improvement in progression-free survival (PFS), response rate and OS [14]. In the overall study population, progression-free survival (PFS) was 7.4 months with cabozantinib versus 3.9 months with everolimus ($p < 0.001$). Subset analyses of patients with bone metastases show persistence of this PFS benefit (7.4 months with cabozantinib vs. 2.7 months with everolimus) [43]. Cabozantinib was also noted to exert a greater reduction in markers of bone turnover, such as N-telopeptide.

5.2. Brain metastases

As noted previously, brain metastases occur in roughly 9% of patients with mRCC. The central nervous system (CNS) is thought to represent a sanctuary site for RCC; preclinical studies show incomplete penetration with agents such as sunitinib and sorafenib [44]. For front-line therapy, the pivotal phase III study of tamsulosin did permit patients with brain metastases – a distinction from previously published front-line trials of VEGF-directed therapies. In patients with mRCC and poor-risk features, tamsulosin was compared to IFN- α and the combination of both agents. Treatment with tamsulosin monotherapy led to an

improvement in OS as compared to IFN- α . Although subset analyses pertaining to patients with brain metastases are not available, the inclusion of this subpopulation provides some justification for offering tamsulosin front-line.

There have been retrospective reviews assessing the safety and efficacy of VEGF-TKIs in mRCC with brain metastasis. In one single institution study, 65 patients with brain metastases from mRCC were identified. The preponderance (80%) received VEGF-directed therapy front-line with a very limited number of neurologic adverse events. In total, neurologic adverse events were noted in five patients, and included radiation necrosis and brain metastasis hemorrhage.

In the second-line setting, the current clinical debate surrounds use of either cabozantinib or nivolumab. Like cabozantinib, nivolumab was assessed in a phase III comparison against everolimus in patients with prior VEGF-directed therapy [12]. Nivolumab was noted to improve both OS and response rate as compared to everolimus. However, it is worthy to note that the phase III assessment of nivolumab did not include patients with brain metastases, while the phase III assessment of cabozantinib did. Thus, in the second-line setting, cabozantinib may be the preferred approach for patients with CNS disease.

6. Conclusion

An understanding of the biology of RCC metastasis will most surely drive advances in systemic therapy in both the adjuvant and metastatic setting. As noted herein, there are multiple studies that are currently addressing the potential role of VEGF- and mTOR-inhibitors as adjuvant treatment. This is a derivative approach based on the observed activity of these agents in the metastatic setting. Although the

majority of adjuvant studies have yet to report, the emerging data from the S-TRAC and ASSURE studies (both exploring VEGF-TKIs) have produced contrasting results. Cautious interpretation of the data will surely be necessary before these approaches are implemented clinically. The potential interplay of immune cells in generating metastases might prompt speculation that adjuvant PD-1 or PD-L1 inhibition could be feasible as adjuvant treatment. These studies will likely be emerging in the near future.

Once RCC does spread, it is critical to note that varying sites of metastasis may carry distinct implications for prognosis. While patients with bone, liver and brain metastases may carry an inferior prognosis relative to the overall population of mRCC patients, patients with pancreatic metastases may have a superior prognosis. At present, there is insufficient evidence to take a tailored approach for each metastatic site. Subset analyses from phase III studies do afford an opportunity to understand the relative benefit of systemic agents in unique settings (e.g., cabozantinib in bone metastases). Furthermore, review of eligibility criteria from these phase III studies do allow the practicing clinician to understand the applicability of data — for instance, patients with brain metastases were not consistently permitted entry in front-line trials. Prospective studies that assess these unique populations may ultimately provide the most relevant insights.

Many gaps remain in our understanding of RCC metastasis. For instance, although the distinct biology of RCC histologies was acknowledged, we have little sense of how this might affect patterns of metastasis. The varied clinical outcomes for different histological subtypes certainly underscores that each may progress through distinct mechanisms. Furthermore, although much emphasis has been placed on optimizing treatment of disease at certain anatomic sites (e.g., bone and brain metastases), it would be helpful to know if other sites of metastases (e.g., pancreatic metastases, adrenal metastases and so on) progress through distinct mechanisms. A better biological understanding of these clinical circumstances will move us towards the goal of personalizing therapy across these settings.

Conflicts of interest

Dr. Pal receives honoraria from Novartis, Medivation and Astellas Pharma, and receives consulting fees from Pfizer, Novartis, Aveo, Genentech, Exelixis, BMS, Astellas and GSK.

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