Therapeutic Options for Variant Renal Cancer: A True Orphan Disease

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ABSTRACT
Variant or nonclear cell renal cell cancer is a rare disease constituting only ~5% to 8% of the metastatic renal cell cancer population. Pathological criteria for the three main variant subtypes, papillary, chromophobe, and collecting duct, have been specified. Nonetheless, there may be subtypes within these variants, many poorly differentiated tumors cannot be reliably classified, and expertise in recognizing specific subtypes is not widespread. Expression analysis and other molecular techniques are beginning to clarify and standardize the pathological classification scheme. Because these classifications are relatively new and the number of patients with any one subtype is limited, little is known about appropriate therapies for patients with metastatic disease. Retrospective series strongly suggest that immunotherapy is not effective in any nonclear cell subtype. Case reports suggest that cytotoxic chemotherapy used for transitional cell cancers may be helpful in patients with collecting duct cancers. A central registry of patients with variant renal cell cancer should be created in which response to various therapies is recorded. Such a registry could provide support for a more formal multi-institutional study investigating a specific drug or regimen.

INTRODUCTION
Until recently, all adult renal epithelial malignancies were considered one disease labeled renal cell carcinoma. During the last 10 years, however, renal cell carcinoma has been increasingly recognized as several different subtypes that have distinct biological attributes, pathological characteristics, and likely oncogenesis (1, 2). As such, renal cell carcinoma is best thought of as several separate diseases for which unique and specific therapeutics must be developed. To begin to develop specific regimens for each subtype, especially the less common ones, a brief review of their biology and the effectiveness of available therapies must be developed. To begin to develop specific regimens for each subtype, especially the less common ones, a brief review of their biology and the effectiveness of available therapy is in order.

Renal cell carcinoma can be divided into classic clear cell carcinoma, papillary carcinoma, chromophobe carcinoma, and collecting duct carcinoma (3). Despite consensus on broad categories of nonclear cell carcinoma of the kidney, additional subsets for which the relationship is somewhat unclear continue to be described. Papillary cancers have, for example, been subdivided into type 1 and type 2 (4), and some controversy remains regarding whether medullary cancers, which have been described almost exclusively in patients with sickle cell disease or trait, are a variant of collecting duct carcinoma (5). In addition, a number of poorly differentiated tumors exist, often classified as sarcomatoid, in which unique pathological characteristics cannot be recognized. Unfortunately, expertise in recognizing even the standard subtypes varies dramatically. Finally, mixed histologies are not uncommon, and the biological significance of such findings is unclear. More importantly, much of the therapeutic literature for renal cell carcinoma either does not subclassify patients by histologic characteristics or uses histologic classifications that are no longer considered standard.

GENETICS AND BIOLOGY
Classic molecular genetics, cytogenetics, and expression profiling have clarified some of the relationships among the various histologic subtypes. The proposed two papillary subtypes are indeed closely related genetically (4). Similarly, chromophobe tumors are molecularly closely related to benign oncocytomas, whereas collecting duct carcinomas and medullary carcinomas are likely more closely related to transitional cell carcinomas (6, 7). Additional expression profiling studies and the use of specific molecular probes and immunohistochemical markers will likely serve to further standardize and define the various renal cell carcinoma subtypes.

Despite the power of molecular profiling, from a therapeutic standpoint it is most important to identify those molecular alterations that are critical to the malignant phenotype of a particular tumor. To this end, the identification of the initiating genetic alterations, similar to the bcr-ABL translocation in chronic myelogenous leukemia, is critical. The study of familial renal cell carcinoma has provided several clues to potential genes critical for the renal cell carcinoma phenotype. In the classic clear cell carcinomas and as described elsewhere (8), study of the von Hippel-Lindau (VHL) syndrome has proved important. The VHL gene product, which is inactivated in the germ line of VHL syndrome patients and in the clear cell renal cell carcinoma tumor cells (9, 10), targets the hypoxia-inducible factor (HIF) transcription factor for ubiquitin-mediated destruction in a normoxic environment (11–13). In the absence of a functional VHL gene product, HIF remains active and mediates the transcription of, among other factors, vascular endothelial growth factor (VEGF), platelet-derived growth factor, and carbonic anhydrase IX (12, 13). These genes contribute to the known hypervascularity of these tumors and may explain the recently described activity of the anti-VEGF antibody bevacizumab, as well as the VEGF receptor and platelet-derived growth factor.
receptor tyrosine kinase inhibitors BAY 43–9006 and SU11248 in this disease (14–16).

Familial nonclear cell renal cell carcinomas have also been described, and several germ line-mutated genes putatively responsible for the observed phenotypes have been identified (2). For example, familial type I papillary cancers have been characterized by MET oncogene mutations that lead to activation of this growth factor receptor tyrosine kinase (17). As a result, the pathway is activated even in the absence of its natural ligand, hepatic growth factor (also known as scatter factor; ref. 18). Familial type II papillary cancers have been associated with cutaneous and uterine leiomyomas and mutations in fumarase hydratase (19–21). The mechanism behind oncogenesis after mutation of this metabolic enzyme is currently not known. Finally, familial Birt-Hogg-Dubé (BHD) syndrome has been associated with multiple renal tumors (usually of the chromophobe subtype) in addition to skin fibrofolliculomas and pulmonary cysts (22). The responsible tumor suppressor gene, the BHD gene, has been identified recently, and some of the mutations in the familial kindreds have been described (23, 24).

However, in all of these cases, and unlike the role of VHL in clear cell cancers, the role of these genes in sporadic forms of each nonclear cell histologic subtype is less clear. For example, MET mutations are very rare in patients with sporadic papillary cancer (25). Although immunohistochemical expression of c-MET has been described in the sporadic forms of papillary renal cell carcinoma, the physiologic relevance of this finding remains unclear (26). The BHD gene is occasionally inactivated in spontaneous renal cell carcinomas of various subtypes, but once again whether this is important to the malignant phenotype remains unknown (27). Perhaps more important is the fact that the role of other members of genetic and metabolic pathways inactivated in the familial nonclear cell cancers have not been investigated. Therefore, unlike the case in clear cell renal cell carcinoma in which VEGF receptor and platelet-derived growth factor receptor can be therapeutically targeted, putative therapeutic targets in nonclear cell renal cell carcinoma remain to be identified.

INCIDENCE AND NATURAL HISTORY

The incidence of nonclear cell carcinoma of the kidney has been reported as ~20% to 30%, with papillary cancers constituting 10% to 15%, chromophobe constituting 5%, and collecting duct constituting ~1% of cases (1, 28). These numbers, however, reflect data from surgical series of primary nephrectomies. Because primary nonclear cell carcinomas tend to have a better prognosis than clear cell carcinomas, the percentage of metastatic nonclear cell carcinomas is much lower (28). Recent reviews from Memorial Sloan-Kettering Cancer Center (New York, NY; ref. 29), the Cytokine Working Group, (30), and the University of Chicago (Chicago, IL; ref. 31) suggest that only 5% to 8% of patients treated in clinical trials have metastatic renal cell carcinoma. Expression profiling, cell lines, and animal models can be attempted. However, only one or two variant cancers are likely to be enrolled in any typically sized Phase II trial, and subgroup analyses could then be performed. If a registry reveals variant histologic subtype. There are, thus, <600 patients who develop any one of these diseases on an annual basis in the United States. With the current rate of enrollment into clinical trials in the adult oncology community, even a cooperative group trial would likely take several years to accrue. This raises the issue of how to wisely choose agents for additional development.

Choosing appropriate agents will clearly depend on improvements in understanding the biology of nonclear cell carcinomas. Expression profiling, cell lines, and animal models can all provide potential predictive information. It is important to recognize, however, that activation or overexpression of a particular pathway is not enough to make that pathway a good therapeutic target. The molecular abnormality has to be necessary for malignant growth.

In the meantime, decisions will have to be based on clinical observations. To this end, a clinical registry of nonclear cell carcinomas may be helpful. Major centers all evaluate two to eight patients with nonclear cell renal cell carcinoma annually and typically treat them with a variety of regimens, including off-protocol therapy, Phase I clinical trials, and occasionally Phase II clinical trials with broad eligibility. If a registry reveals
that several patients with a specific renal cell carcinoma subtype have responded to a particular class of agents, this might be sufficient to justify a larger cooperative group clinical trial. The value of such a registry can be illustrated by anecdotal observation of responses to Gleevec in gastrointestinal stromal tumors, which led to identification of the pathognomonic c-kit inactivation in these tumors and a collaborative group of investigators dedicated to performing therapeutic studies (34).

CONCLUSIONS
Nonclear cell or variant metastatic renal cell carcinomas are unique diseases that will require unique therapeutic approaches. The recognition of these subtypes by general pathologists is improving, and molecular diagnostics are likely to additionally improve the diagnostic accuracy. Nevertheless, the rarity of these diseases in the metastatic setting means that they are truly orphan diseases for which it is extremely difficult to conduct clinical trials. Therefore, a clinical registry should be created in which response to various off-protocol and early phase clinical trials are recorded. In addition to ongoing work on the biological pathways important in these diseases, such a registry may provide clues to which classes of agents may be therapeutically effective.

OPEN DISCUSSION

Dr. Michael B. Atkins: Do you know of any patients with nonclear cell histology who have had a response to some of the other targeted agents?

Dr. Daniel J. George: We had two patients with chromophobe tumors who had a minor response on PTK787. I think that is all you are going to find in these kinds of studies. These tumors are not that dissimilar in terms of their biology, although genetically they are. I think there is a reason to try some of these VEGF targeted strategies in patients with nonclear cell histology.

Dr. Atkins: But the reason is the same reason you would try it in colon cancer or in any other cancer.

Dr. George: Yes, that’s right. If I have a patient with a chromophobe tumor, I would be willing to enroll him on a Phase I study with a VEGF inhibitor.

Dr. William G. Kaelin, Jr.: Would it be fair to make the argument that there is as much heterogeneity with what we call clear cell carcinoma as with other types of cancer?

Dr. Walter M. Stadler: There is a huge difference between clear cell and other cell types, but within the clear cell, there are different subgroups. We don’t know why, but that is the reason for the difference in response within that larger clear cell group.

Dr. Kaelin: Do we decide that from now on we are going to treat the VHL mutant clear cell population differently as opposed to treating all clear cell the same or do we want to allow for empiricism and include everybody?

Dr. Bin Tean Teh: For this case, that is, the so-called nonclear cell, it is very important to make sure that the diagnosis is correct.

Dr. Stadler: We need better diagnostics, but I think we are fairly close to being very accurate in terms of how we diagnose these different subtypes. It still leaves us with the question of how we develop therapies for some of these subtypes that are extremely rare. We have too few patients, too many drugs, and not necessarily a good rationale.

Dr. Robert A. Figlin: Some of my most difficult decisions these days are deciding what to do with the patients with pure papillary tumors. There are a variety of things out there, but there is not a lot of information that suggests that we have any knowledgeable approach to this patient population. Are we talking about papillary tumors or are we talking about tumors that have wild-type VHL?

Dr. Stadler: We are talking about the nonclassic clear cells, those with wild-type VHL.

Dr. Atkins: So, this could be up to 40% of the patients.

Dr. Figlin: Correct. I’m just wondering whether we need to start thinking about approaching kidney cancer as diseases with either mutated or nonmutated VHL.

REFERENCES


34. Verweij J, van Oosterom A, Blay JY et al. Imatinib mesylate (STI-571 Gleevec, Gleevec) is an active agent for gastrointestinal stromal tumours, but does not yield responses in other soft-tissue sarcomas that are unselected for a molecular target. Results from an EORTC Soft Tissue and Bone Sarcoma Group phase II study. Eur J Cancer 2003;39:2006–11.
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