Review Article

Progress of Molecular Targeted Therapies for Advanced Renal Cell Carcinoma

Alessandro Conti, Matteo Santoni, Consuelo Amantini, Luciano Burattini, Rossana Berardi, Giorgio Santoni, Stefano Cascinu, and Giovanni Muzzonigro

1 Department of Clinical and Specialist Sciences, Urology, Polytechnic University of the Marche Region, AOU Ospedali Riuniti Umberto I-GM Lancisi and G Salesi, Ancona, Italy
2 Medical Oncology, Polytechnic University of the Marche Region, AOU Ospedali Riuniti Umberto I-GM Lancisi and G Salesi, Ancona, Italy
3 School of Pharmacy, Section of Experimental Medicine, University of Camerino, Italy

Correspondence should be addressed to Alessandro Conti; alessandro.conti@hotmail.com

Received 30 April 2013; Revised 13 July 2013; Accepted 3 August 2013

Copyright © 2013 Alessandro Conti et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Vascular endothelial growth factor (VEGF) plays a crucial role in tumor angiogenesis. VEGF expression in metastatic renal cell carcinoma (mRCC) is mostly regulated by hypoxia, predominantly via the hypoxia-induced factor (HIF)/Von Hippel-Lindau (VHL) pathway. Advances in our knowledge of VEGF role in tumor angiogenesis, growth, and progression have permitted development of new approaches for the treatment of mRCC, including several agents targeting VEGF and VEGF receptors: tyrosine kinase pathway, serine/threonine kinases, α5β1-integrin, deacetylase, CD70, mammalian target of rapamycin (mTOR), AKT, and phosphatidylinositol 3-kinase (PI3K). Starting from sorafenib and sunitinib, several targeted therapies have been approved for mRCC treatment, with a long list of agents in course of evaluation, such as tivozanib, cediranib, and VEGF-Trap. Here we illustrate the main steps of tumor angiogenesis process, defining the pertinent therapeutic targets and the efficacy and toxicity profiles of these new promising agents.

1. Introduction

The development of targeted agents has completely changed the therapeutic landscape of metastatic renal cell carcinoma (mRCC). Starting from sorafenib in 2005, several agents have been sequentially approved by the US Food and Drug Administration (FDA), such as sunitinib, bevacizumab plus interferon, everolimus, temsirolimus, pazopanib, and at last axitinib. Nevertheless, despite recent success, complete responses to antiangiogenic therapies are rare, and patients do progress in response to antivascular endothelial growth factor (VEGF) targeted therapies. The increasing awareness of the molecular mechanisms underlying tumor angiogenesis in RCC parallels the development of novel antiangiogenic agents, mainly directed against VEGF-A, platelet-derived growth factor (PDGF) receptors (PDGFRs), and VEGF receptor-2 (VEGFR-2), with a list of novel targets currently in course of evaluation.

In RCC, the loss of 3p induces a high incidence of Von Hippel-Lindau (VHL) tumor suppressor gene inactivation, which promotes hypoxia-inducible factor-1 (HIF-1) dysregulation [1], and leads to increased expression of VEGF and other proangiogenic factors [2]. HIF-1 activity is essential for genetic stability, proliferation, migration, and survival, contributing also to tumor cell metabolism and drug and radiation resistance [3, 4].

Tumor angiogenesis is the result of the interaction among several components of tumor microenvironment. This multistep process involves the migration and proliferation of endothelial cells (ECs), the recruitment of hematopoietic progenitor cells (HPCs), highly proliferative circulating endothelial cells (CECs), and endothelial progenitor cells (EPCs), the detachment and migration of pericytes, and the cooption of neighbouring preexisting vessels [5]. Several signaling pathways are involved in tumor angiogenesis, such as VEGF,
which plays a crucial role in RCC carcinogenesis [6, 7], PDGF, the mammalian target of rapamycin (mTOR), VHL, a tumor suppressor gene, and HIF (Figure 1). The transit from an anti- to a proangiogenic phase, also called angiogenic switch, leads to the rapid formation of blood vessels [8]. The insufficient vascularisation accompanying rapid tumor growth leads to hypoxia [9, 10].

VEGF promotes EC migration and proliferation, thus stimulating extracellular matrix (ECM) and EC basement membrane digestion by plasminogen activator (PA) and metalloproteinases (MMPs). Tumor vessel sprouting requires, at first, the removal of pericytes from preexisting blood vessels. As a result of this process, ECs begin to proliferate and migrate to form new vessels, subsequently surrounded by mesenchymal cells differentiated into pericytes. Tumor cell-derived VEGF induces also the recruitment of CECs and EPCs. CECs are generally accepted as cells expressing the von Willebrand factor (vWF), CD146, CD34, and VE-cadherin. Their clinical significance is still under investigation. Recent studies have observed increased levels of CECs in RCC patients treated with sunitinib, likely caused by the drug targeting of immature tumor vessels [11]. EPCs express CD133, CD34, and VEGFR-2 [12]. Circulating EPCs begin to differentiate by losing the CD133 marker and gaining EC-specific markers such as vWF, CD31, and VE-cadherin [13], thus enhancing tumor development and growth [14]. In 2003, Rafii et al. observed that MMP-9 was capable of mobilizing EPCs [15]. Their migration is mediated by the chemokine stromal cell-derived factor-1 (SDF-1), which is upregulated during hypoxia [16].

A related circulating cell population is the HPC, expressing VEGFRI. Along with EPCs expressing VGRF2, they colonize the sprouting vessel connective tissue and also provide growth factors and cytokines to create a microenvironment favouring the start of metastatic process, in which tumor cells can localize and proliferate [17]. Recently, Powles et al. have demonstrated the association between increased numbers of circulating HPCs and poor outcome in patients with mRCC treated with sunitinib [18].

The complete formation of a lumen in novel sprouting vessels needs the contribution of stalk cells, stimulated by VEGF, ECM, and EC-derived EGF-like domain 7 (Egf7), which promotes the separation and proper spatial arrangement of the angioblasts and allows subsequent assembly of vascular tubes [19].

The detachment of pericytes during tumor angiogenesis is mediated by VEGF and angiopoietin-2 (Ang-2)/type I tyrosine kinase receptor 2 (TIE 2). The moved pericytes are recruited by PDGF expressed in ECs, surrounding and stabilizing vessel sprouts [20].

The expression of angiogenic factors in RCC is also regulated by VHL-independent signaling pathways, such as the oncoprotein homologue of the mouse double minute 2 (HDM2) that leads to the constitutive expression of HIFα. HDM2 also regulates the protein levels of HIF angiogenic targets, such as VEGF, PA inhibitor-1 (PAI-1), and endothelin-1 (ET-1) [21].

Therefore, in this review we illustrate the progress of molecular targeted therapy, describing the emerging agents in course of evaluation and the future directions in the treatment of RCC patients.

2. Tyrosine Kinase Inhibitors

“Enlarged” Family

In the last two years, the list of novel anti-angiogenic agents under evaluation has progressively enlarged, including now axitinib, tivozanib, dovitinib, cediranib, linifanib, volociximab, regorafenib, and other new emerging molecules [22]. In January 2012, axitinib was approved in the second line treatment of mRCC patients. Axitinib is an oral inhibitor of VEGFR, PDGFR, and c-kit, and it may afford a relevant contribution for the future management of RCC patients. In a phase II trial enrolling 62 treatment-refractory RCC patients progressed on sorafenib and treated with oral 5 mg axitinib twice daily, the median progression-free survival (PFS) was 7.4 months. Grades 3 to 4 adverse events included hand-foot syndrome (16.1%), fatigue (16.1%), hypertension (16.1%),
dyspnea (14.5%), diarrhea (14.5%), dehydration (8.1%), and hypotension (6.5%) [23]. In addition, another phase II trial with axitinib, enrolling cytokine-refractory nephrectomised patients, showed response rate (RR) of 44.2% and a median time to progression (TTP) of 15.7 months [24]. The AXIS phase III trial, comparing the effectiveness of axitinib and sorafenib in pretreated advanced RCC patients, has shown a significant median PFS (6.7 versus 4.7 months), with a more pronounced benefit for patients after immunotherapy (12.1 versus 6.5 months) [25]. With regard to the front-line use of axitinib, the preliminary results shown at the 2012 annual meeting of the American Society of Clinical Oncology (ASCO) from the randomized phase II study AGILE 1046 demonstrated a median PFS of more than 1 year and an overall response rate of 40.2% for axitinib [26]. Furthermore, several trials are presently ongoing to evaluate the role of axitinib as neoadjuvant therapy (NCT01263569), in combination with everolimus (NCT01334073), or coadministered with PF-04856884, a selective angiopoietin (Ang)-2 inhibitor (NCT01441414).

Among the emerging agents, tivozanib (AV-951) has been demonstrated to be effective in the first line setting of mRCC patients with prior nephrectomy [27]. Tivozanib is a potent inhibitor of VEGFR-1, 2, and 3, cKit, and PDGFR. Hypertension (50.0%) and dysphonia (21.7%) were the most commonly reported treatment-related adverse events of any grade. There was a low incidence of diarrhea (12.1%), fatigue (8.1%), stomatitis (4.4%), and hand-foot syndrome (3.7%). In the phase III “TIVO-1” study [28], a total of 517 patients were randomized to tivozanib (N = 260) or sorafenib (N = 257) in first line advanced RCC patients. Tivozanib showed a statistically significant improvement in median PFS compared to sorafenib (11.9 versus 9.1 months) in the subpopulation of patients who were pretreated with systemic therapy including cytokines. In patients who were treatment naive (70% of total study population), tivozanib showed a statistically significant improvement in PFS, with a median PFS of 12.7 months compared with 9.1 months for sorafenib (HR 0.756, 95% CI 0.580–0.985; P = 0.037). Tivozanib demonstrated favorable tolerability, with a lower rate of dose interruptions (18% versus 35%, P < 0.001) and reductions (14% versus 44%, P < 0.001). The most common ≥ grade 3 adverse events (AEs) due to tivozanib compared to sorafenib were hypertension (25% versus 17%), hand-foot syndrome (2% versus 17%), diarrhea (2% versus 6%), fatigue (5% versus 4%), and neutropenia (2% versus 2%). While the progression-free survival was improved, the overall survival (OS) showed a trend toward a detrimental effect with the tivozanib arm with a median OS of 28.8 months versus 29.3 months in the sorafenib arm based on the pre-new drug application (NDA) meeting with the US Food and Drug Administration (FDA) [29] which later led to the FDA OADAC meeting to disapprove tivozanib as an indication for RCC.

A phase I study has been completed to evaluate the safety of tivozanib in combination with temsirolimus in subjects with mRCC (NCT00563147).

With regard to the third line treatment of mRCC patients, dovitinib seems to represent a valid option. It is a fibroblast growth factor receptor (FGFR) and VEGFR inhibitor, presently in course of evaluation in a phase III trial (NCT01223027). The most common adverse events shown in the phase II study were nausea (80%; G3/5%), diarrhea (70%), vomiting (65%), asthenia (50%; G3/5%), anorexia (45%; G3/5%), headache (30%; G3/5%), hypertension (25%; G4/5%), and rash (23%; G3/5%). In a phase II trial enrolling 59 previously treated patients, dovitinib was administered with a dose schedule of 500 mg/day 5 days on/2 days off. In this study, PFS and OS were 6.1 and 16 months, respectively [30]. Results are awaited from a phase III trial (NCT01223027) enrolling 550 patients who must have received one VEGF-targeted therapy and one prior mTOR inhibitor therapy to evaluate dovitinib versus sorafenib in the third line setting of mRCC treatment.

Recent advances in understanding the role of fibroblast growth factor factor 2 (FGF2) and FGF receptor (FGFR) in modulating resistance to sunitinib [31] led to the development of PD173074, a reversible FGFR and VEGFR inhibitor. Thus, FGF2 supports endothelial proliferation and de novo tubule formation in the presence of sunitinib, suppressing sunitinib-induced retraction of tubules. Currently, several studies are analyzing the efficacy and safety of PD173074 in small cell lung cancer and RCC.

At this time, the list of emerging TKIs under study in phase II trials includes cediranib, linifanib, regorafenib, brivanib, vandetanib, lenvatinib, and several other agents. Cediranib (AZD2171) is an oral inhibitor of VEGFR-3, PDGFRβ, and c-kit. In 2012, Mulders et al. have published the results from a phase II trial (NCT00423332) in 71 previously untreated mRCC patients randomized to receive cediranib (n = 53) or placebo (n = 18). They revealed 34% PR and 47% stable disease (SD), and cediranib was generally well tolerated [32]. Furthermore, another phase II trial (COSAK) is ongoing to assess the efficacy of cediranib 30 mg versus cediranib 30 mg plus 175 mg saracatinib (AZD0530), an Src Family oral inhibitor, in patients with relapsed metastatic clear cell RCC (ccRCC).

Linifanib (ABT-869) is a potent inhibitor of VEGFR, PDGFR, fms-like tyrosine kinase 3 (FLT3), c-kit, and colony stimulating factor-1 receptor (CSFIR). In 2012, Tannir et al. have published their results [33] from an open-label multicenter trial (NCT00486538) in 53 patients previously treated with sunitinib, receiving oral linifanib 0.25 mg/kg (12.5–25.0 mg) daily. They showed 13.2% overall RR, with a median PFS and OS of 5.4 and 14.5 months, respectively.

Regorafenib (BAY 73-4506) is an orally multikinase inhibitor targeting VEGFR, c-kit, RET, FGFR, PDGFR, and serine/threonine kinases (RAF and p38MAPK). A phase II trial (NCT00664326) on 33 patients treated with BAY 73-4506 160 mg once daily on a 3-week on/1-week off schedule showed 27% PR and a 42% SD [34].

Brivanib and vandetanib represent two more members of the VEGF-related antiangiogenic family. Brivanib is an oral, dual VEGFR-2 and FGFR-1 tyrosine kinases inhibitor. A phase II, open-label investigation conducted to assess is activity in mRCC patients has been opened in November 2010 (NCT01253668). On the other hand, vandetanib, also known as ZD6474, is an antagonist of VEGF and EGFR. A phase
II trial (NCT01372813) has been terminated for insufficient accrual.

In 2006, Jermann et al. [35] published the results of a phase II trial of gefitinib, a low-molecular-weight epidermal growth factor receptor (EGFR) TKI, in patients with locally advanced, metastatic, or relapsed RCC. They did not observe objective responses (OR), although 14 patients (53.8%) had SD. Treatment was generally well tolerated. Furthermore, gefitinib was also evaluated in combination with pegylated IFNα, without obtaining significant results [36].

The development of XL999 and tandutinib was stopped due to their toxicity profiles. In 2006, a phase II trial (NCT00277316) was opened to investigate the activity of XL999, a multiple inhibitor of VEGFR, PDGFR, FGFR, FLT-3, and Src. The employment of this molecule was stopped due to the cardiac AEs revealed in the participants of this study. In the same year, Shepard et al. [37] evaluated the efficacy and tolerability of tandutinib, an oral inhibitor of type III tyrosine kinase receptor kinases, observing absence of clinical activity and excessive toxicity in these patients.

Finally, the list of novel TKI agents presently under study also includes crizotinib, an anaplastic lymphoma kinase (ALK), and c-Met TKI (CREATE, NCT01524926) and BIBF 120, which is under evaluation versus sunitinib in untreated mRCC patients (NCT01024920).

3. Anti-VEGFR Dual Strategy: mAb and Derivatives

Since the approval of bevacizumab in 2009 for the treatment of RCC patients, several VEGF/VEGFR blocking agents have been developed, such as VEGF-Trap and ramucirumab. VEGF-Trap (AVE0005, ziv-aflibercept) is a VEGF-blocking agent with both higher affinity and activity than bevacizumab [38]. It is a derivative of VEGFR1 and VEGFR2 [38], with minimal interactions with extracellular matrix, and this property may be the key to its satisfying pharmacokinetic profile. At present, a phase II trial (NCT00357760) is in course in patients who must have received at least one prior treatment with a TKI for metastatic ccRCC. VEGF-Trap (ziv-aflibercept). The main side effects revealed in the phase I trial were proteinuria (37%), fatigue (32%), injection site reactions (18%), nausea (17%), myalgia and anorexia (16% each), hypertension (13%), and voice hoarseness (11%), with 7% of grades 3-4 events [39].

Ramucirumab (IMC-I12-IB) is a fully human IgG1 mAb targeting VEGFR-2, used in a phase II trial (NCT00515697). The results obtained, however, (49% SD lasting over 5 months, 6 months PFS) must be confirmed in randomized trials on TKI-refractory mRCC patients.

4. Targeting mTOR Pathway at Several Levels: mTOR, AKT, and PI3K Inhibitors

Recent strategies against mTOR pathway include the use of selective inhibitors of mTOR, AKT, or PI3K and dual PI3K/mTOR inhibitors. Between the emerging mTOR inhibitors, ridaforolimus (AP23573) [40] and AZD8055 [40] have been demonstrated to be active in RCC patients. Ridaforolimus is an mTOR complex 1 (MTORC1) selective inhibitor that has been studied in phase I trial [39] administered as a 30-minute intravenous infusion once daily for 5 consecutive days every 2 weeks (QDx5) in a 28-day cycle. On the basis of this study, a dose of 12.5 mg/d is being evaluated in phase II trials. Ridaforolimus is currently under evaluation even in combination with vorinostat, an HDAC inhibitor (NCT01169532), and with AKT inhibitor MK2206 or γ-secretase inhibitor MK-0752 in patients with advanced solid tumors (NCT01295632).

On the other hand, AZD8055, a dual mTORC1 and mTORC2 inhibitor, has been tested in a phase I clinical trial at different doses (NCT00731263), showing a maximum tolerated dose (MTD) of 90 mg BID [41].

Moreover, perifosine, an AKT and MAP kinase inhibitor, has been evaluated in a phase II trial (NCT00448721) in 24 patients who had experienced disease progression after receiving either sorafenib or sunitinib. This study revealed 8% PR, 42% SD at 12 weeks, and a median PFS of at 19 weeks. The most common grades 3 and 4 adverse events were dyspnea (8%), hynonatremia (8%), pulmonary embolism (4%), and arthralgia (4%) [42]. Two separate phase I trials have investigated its use in combination with sunitinib (NCT00399152) or sorafenib (NCT00398814), observing in both cases a good tolerability and promising results in stabilizing kidney cancer. The efficacy of MK2206, another AKT inhibitor, is presently under evaluation in a phase II trial (NCT01239342) versus everolimus in refractory RCC patients.

Concerning PI3K inhibitors, BKI-120 is presently under evaluation in several clinical trials, administered alone or in combination with bevacizumab in patients who have failed at last one prior anti-VEGF therapy (NCT01283048).

Finally, this family also includes dual PI3K/mTOR inhibitors, such as NVP-BEZ235 and GDC-0980. NVP-BEZ235, a dual PI3K/mTOR inhibitor, has been demonstrated to reduce in vitro cell proliferation by inducing nuclear translocation of p27 and to downregulate Akt, Mnk-1, eIF4E, and 4EBP-1 phosphorylation and cyclin D1 and HIF2α compared to rapamycin [43]. In July 2011, Roulin et al. evaluated in vitro the combination between NVP-BEZ235 and sorafenib, resulting in reduced tumour cell proliferation and increased tumour cell apoptosis in vitro [44]. Moreover, NVP-BEZ235 is under study with everolimus in a phase I trial (NCT01482156). With regard to GDC-0980, it is under evaluation in comparison with everolimus in mRCC patients progressed on VEGF-targeted therapy (NCT01442090).

5. Novel Emerging Targets and Future Directions

The role of angiopoietin/Tie2 pathway in tumor angiogenesis has led to the development of several inhibiting agents, such as AMG 386 and PF-4856884. Their action alone or in combination with TKIs is subject of ongoing clinical trials. The efficacy and tolerability of trebananib (AMG 386) plus sorafenib have been tested in 152 ccRCC patients receiving sorafenib 400 mg orally twice daily plus intravenous AMG 386 at 10 mg/kg (arm A) or 3 mg/kg (arm B) or placebo (arm C).
recently, the chimeric mAb cG250 has been demonstrated in in vitro studies to induce antibody-dependent cellular cytotoxicity (ADCC) against G250+ RCC cells [51]. The employment of G250 in the diagnosis and treatment of RCC is under evaluation.

Histone deacetylase (HDAC) determines the acetylation status of histones, thus regulating gene expression. In 2004, Shao et al. reported that aberrant HDAC activity plays a relevant role in carcinogenesis [52]. HDAC catalyses the removal of acetyl groups from N-acetyl lysine on a histone. The effects of HDAC inhibitors include the activation of cyclin-dependent kinase inhibitor p21WAF1/CIP1 [53] and the p53/p21 pathway activation [54]. The mechanism underlying HDAC-related apoptosis in tumor cells is still unclear, but it seems to involve the recruitment of Bcl-2 family members. LBH589 (panobinostat) is a nonselective HDAC inhibitor interfering with the deacetylation process. Its activity has been tested in RCC and haematological malignancies. Results from a phase II study [55] conducted in twenty patients who had received at least one prior TKI and one mTOR inhibitor, demonstrated no ORs, although panobinostat was well tolerated. Recently, a phase II trial (NCT01037257) has been opened to investigate its activity in combination with everolimus.

In the last years, the list of therapeutic targets has widely enlarged, including at present the signal transducer and activator of transcription 3 (STAT 3), Notch homolog 1 translocation-associated (Notch-1), dysmorphic ECs, and nucleolin. Preliminary results from further clinical trials are helpful to assess their role in the future RCC treatment landscape.

STAT 3 is involved in several signaling pathways, regulating cell survival, proliferation, and angiogenesis, and it is aberrantly activated in RCC [56]. In a recent study, the activity of WPI066, a STAT 3 inhibitor, has been tested in vitro in RCC lines and in vivo on murine xenografts; the results obtained show an antiproliferative activity related to a reduction in tubulogenesis [57].

Notch pathway is involved in cell communication and angiogenesis representing a potential target for anticancer therapy. The development of gamma-secretase/Notch signaling pathway inhibitor RO4929097 has led to novel perspectives, and it has been approached in a phase I trial (NCT0114569) in mRCC patients that have failed VEGF/VEGFR therapy.

The main objective of vascular targeting agents is to inhibit tumor angiogenesis and to avoid the formation of new sprouting vessels. A different approach may be realized by the use of vascular-disrupting agents (VDAs). VDAs recognize and disrupt the already existing tumor vessels by targeting dysmorphic ECs. VDAs include small molecules and ligand-directed agents. Most of the small molecules are tubulin inhibitors, while ligand-directed agents are mainly flavonoids. Their action results in a local production of TNF-alpha and other cytokines.

Finally, the discovery and development of anticancer aptamers may provide an important contribution to RCC treatment. Aptamers are short DNA, RNA, or peptide oligomers able to assume a specific and stable three-dimensional shape in vivo [58]. The 26-mer DNA aptamer AS1411 is currently undergoing clinical evaluation in acute myeloid leukemia (AML), RCC (NCT00740441), and several other solid tumors. AS1411 is internalized into tumor cells by specific binding to nucleolin. Nucleolin is involved in several cellular activities, such as mRNA stabilization and ribosome assembly [59]. AS1411 activity seems to be related to its decoy action, which interferes with the stabilization of bcl-2 mRNA by nucleolin, inducing cell death [60].
Table 1: Novel targeted agents currently under evaluation for mRCC.

| Agent        | Description                                                                 | Trial ID number | Phase | Design                                                                 |
|--------------|------------------------------------------------------------------------------|-----------------|-------|------------------------------------------------------------------------|
| Brivanib     | Dual VEGFR2 and FGFR-1                                                        | NCT01253668     | II    | RCC patients after prior treatment with TKI or bevacizumab            |
| Crizotinib   | Alk and c-MET TKI                                                            | NCT01524926     | II    | Patients with solid tumors                                            |
| BIBF 120     | VEGFR 1–3 PDGFR and FGFR TKI                                                 | NCT01024920     | II    | versus sunitinib in untreated mRCC patients                           |
| VEGF-Trap    | Soluble decoy receptor; derivative of VEGFRI                                 | NCT00357760     | II    | ccRCC patients after at least 1 prior treatment with TKI              |
| Ridaforolimus| MTORC1 selective inhibitor                                                    | NCT01169532     | I     | In combination with vorinostat in patients with solid tumors         |
| MK-2206      | AKT inhibitor                                                                 | NCT01295632     | I     | In combination with MK2206 or γ-secretase inhibitor MK-0752 in patients with advanced solid tumors |
| NVP-BEZ235   | Dual PI3K/mTOR inhibitor                                                      | NCT01482156     | I     | Versus everolimus in refractory RCC patients                          |
| GDC-0980     | Dual PI3K/mTOR inhibitor                                                      | NCT01442090     | II    | In competition with everolimus in patients with advanced solid tumors |
| AMG-386      | Ang-1/2 inhibitor                                                             | NCT01548482     | II    | In combination with temsirolimus in patients with advanced solid tumors |
| MDX-1203     | Anti-CD70 Ab-drug conjugate                                                   | NCT00944905     | I     | Pretreated ccRCC or B-cell non-Hodgkin's lymphoma                      |
| MDX-1411     | Anti-CD70 Ab-drug conjugate                                                   | NCT00656734     | I     | ccRCC pts treated with up to 6 prior systemic therapies               |
| SGN-75       | Anti-CD70 Ab-drug conjugate                                                   | NCT01015911     | I     | Pretreated ccRCC or B-cell non-Hodgkin’s lymphoma                     |
| Girentuximab | Chimeric mAb cG250                                                           | NCT00087022     | III   | Adjuvant cG250 versus placebo in pts with ccRCC and high risk of recurrence |
| cG250-Lut177 | Lutetium-177 labeled cG250                                                    | NCT00142415     | II    | pts with advanced and progressive ccRCC                               |
| 90Y-cG250    | Yttrium-90 labeled cG250                                                      | NCT00199875     | I     | pts with advanced and progressive ccRCC                               |
| Panitumumab  | Anti-EGFR mAb                                                                 | NCT00425035     | II    | mRCC pts naive or after cytokine treatment                            |
| Vorinostat   | HDAC inhibitor                                                                | NCT00278395     | II    | mRCC pts naive or after cytokine treatment                            |
| RO4929097    | γ-secretase/Notch inhibitor                                                   | NCT01141569     | II    | ccRCC pts after anti-VEGF and/or mTOR inhibitor and/or immunotherapy failure |
| ASI411       | 26-mer DNA aptamer                                                            | NCT00740441     | II    | ccRCC pts after at least 1 prior treatment with TKI                   |

6. Discussion

The advent of targeted therapy has dramatically improved the outcome of mRCC patients. From the approval of sorafenib in 2005, the FDA sequentially approved sunitinib in 2006, temsirolimus in 2007, everolimus, bevacizumab combined with interferon alpha (IFN-α), and pazopanib in 2009, and lastly axitinib in 2012. Based on the results obtained by these agents, a large number of drugs have been developed and are currently under evaluation in RCC patients (Table 1). Nevertheless, the efficacy of these agents seems to be influenced by several factors. The analysis of clinical trials described in this review reveals that toxicity and drug resistance are the main driving forces in continuing research and development of novel agents. The toxicity profiles of emerging molecules result mild to moderate, showing relevant grade 3 and grade 4 adverse event rates (range 7–16.1%). Only volociximab showed no serious adverse events. Although there were differences between the frequencies of PR observed in these trials, all investigated agents were characterized by the lack of CR. These negative data should be a strong stimulus for the creation of new therapeutic approaches.

Primary or acquired resistance to TKIs and mTOR inhibitors has become a major focus for cancer researchers. Primary resistance is less common and seems to be linked to an intrinsic redundancy of tumor available angiogenic signals. On the other hand, acquired resistance is the expression of an “angiogenic switch,” which consists in upregulation of the existing VEGF pathway associated with the recruitment of alternative proangiogenic factors. Current designed preclinical and clinical trials include sequential and combination regimens, aimed at providing a complete blockade of the VEGF pathway. Observing preliminary results, in spite of a mild improvement of patients’ outcome, combined therapies have shown a moderate to severe increase of adverse events. Beyond the toxicity reported, the realization of combined regimens is also limited by their high costs that make them feasible in experimental trials but not in clinical practice.
Moreover, the development of novel effective agents parallels the request of new clinical and molecular predictive and prognostic biomarkers. Their role should be even more relevant if associated with an improved evaluation of tumoral response to antiangiogenic therapy. Finally, an extended knowledge of tumor carcinogenesis process may afford novel tools to optimize treatment regimens. Further clinical trials are still needed to improve the outcome of mRCC patients.

7. Conclusion

Data gathered from ongoing trials will surely improve the management of mRCC patients, even if it is difficult to define the relevance of each one’s individual contribution. Preliminary results from ongoing trials thus constitute a basis for moderate enthusiasm, but a dramatic improvement of mRCC patient outcomes seems still so far.

Conflict of Interests

The authors declare no conflict of interests.

References

[1] M. Zimmer, D. Doucette, N. Siddiqui, and O. Iliopoulos, “Inhibition of hypoxia-inducible factor is sufficient for growth suppression of VHL−/− tumors,” Molecular Cancer Research, vol. 2, no. 2, pp. 89–95, 2004.
[2] B. I. Rini and E. J. Small, “Biology and clinical development of vascular endothelial growth factor-targeted therapy in renal cell carcinoma,” Journal of Clinical Oncology, vol. 23, no. 5, pp. 1028–1043, 2005.
[3] G. L. Semenza, “Targeting HIF-1 for cancer therapy,” Nature Reviews Cancer, vol. 3, no. 10, pp. 721–732, 2003.
[4] K. Podar and K. C. Anderson, “A therapeutic role for targeting c-Myc/Hif-1-dependent signaling pathways,” Cell Cycle, vol. 9, no. 9, pp. 1722–1728, 2010.
[5] J. Holash, P. C. Maisonpierre, D. Compton et al., “Vessel cooption, regression, and growth in tumors mediated by angiopoietins and VEGF,” Science, vol. 284, no. 5422, pp. 1994–1998, 1999.
[6] B. Hemmerlein, A. Kugler, R. Özisik, R.-H. Ringert, H.-J. Radzun, and P. Thelen, “Vascular endothelial growth factor expression, angiogenesis, and necrosis in renal cell carcinomas,” Virchows Archiv, vol. 439, no. 5, pp. 645–652, 2001.
[7] N. Tsuchiya, K. Sato, T. Akae et al., “Quantitative analysis of gene expressions of vascular endothelial growth factor-related factors and their receptors in renal cell carcinoma,” Tohoku Journal of Experimental Medicine, vol. 195, no. 2, pp. 101–113, 2001.
[8] G. Bergers and L. E. Benjamin, “Tumorigenesis and the angiogenic switch,” Nature Reviews Cancer, vol. 3, no. 6, pp. 401–410, 2003.
[9] X. Na, G. Wu, C. K. Ryan, S. R. Schoen, P. A. Di Santagne, and E. M. Messing, “Overproduction of vascular endothelial growth factor related to von Hippel-Lindau tumor suppressor gene mutations and hypoxia-inducible factor-1α expression in renal cell carcinomas,” Journal of Urology, vol. 170, no. 2 I, pp. 588–592, 2003.
[10] N. S. Levy, S. Chung, H. Furneaux, and A. P. Levy, “Hypoxic stabilization of vascular endothelial growth factor mRNA by the RNA-binding protein HuR,” The Journal of Biological Chemistry, vol. 273, no. 11, pp. 6417–6423, 1998.
[11] L. Vroeling, A. A. M. Van Der Veldt, R. R. De Haas et al., “Increased numbers of small circulating endothelial cells in renal cell cancer patients treated with sunitinib,” Angiogenesis, vol. 12, no. 1, pp. 69–79, 2009.
[12] S. M. Bauer, R. J. Bauer, and O. C. Velazquez, “Angiogenesis, vasculogenesis, and induction of healing in chronic wounds,” Vascular and Endovascular Surgery, vol. 39, no. 4, pp. 293–306, 2005.
[13] H. Bompais, J. Chagraoui, X. Cartron et al., “Human endothelial cells derived from circulating progenitors display specific functional properties compared with mature vessel wall endothelial cells,” Blood, vol. 103, no. 7, pp. 2577–2584, 2004.
[14] S. Bruno, B. Bussolati, C. Grange et al., “CD133+ renal progenitor cells contribute to tumor angiogenesis,” American Journal of Pathology, vol. 169, no. 6, pp. 2223–2235, 2006.
[15] S. Rafii, S. Aveicilla, S. Shmelkov et al., “Angiogenic factors reconstitute hematopoiesis by recruiting stem cells from bone marrow microenvironment,” Annals of the New York Academy of Sciences, vol. 996, pp. 49–60, 2003.
[16] D. J. Ceradini, A. R. Kulkarni, M. J. Callaghan et al., “Progenitor cell trafficking is regulated by hypoxic gradients through HIF-1 induction of SDF-1,” Nature Medicine, vol. 10, no. 8, pp. 858–864, 2004.
[17] R. N. Kaplan, R. D. Riba, S. Zacharoulis et al., “VEGFR1-positive haematopoietic bone marrow progenitors initiate the pre-metastatic niche,” Nature, vol. 436, no. 7069, pp. 820–827, 2005.
[18] T. Powles, S. Chowdhury, J. Shamash et al., “Increased haematopoietic progenitor cells are associated with poor outcome in patients with metastatic renal cancer treated with sunitinib,” Annals of Oncology, vol. 22, no. 4, pp. 815–820, 2011.
[19] L. H. Parker, M. Schmidt, S.-W. Jin et al., “The endothelial-cell-derived secreted factor Egfl7 regulates vascular tube formation,” Nature, vol. 428, no. 6984, pp. 754–758, 2004.
[20] C. Betsholtz, P. Lindblom, and H. Gerhardt, “Role of pericytes in vascular morphogenesis,” EXS, no. 94, pp. 115–125, 2005.
[21] V. A. Carroll and M. Ashcroft, “Regulation of angiogenic factors by HDM2 in renal cell carcinoma,” Cancer Research, vol. 68, no. 2, pp. 545–552, 2008.
[22] M. Santoni, M. Rizzo, L. Burattini, R. Berardi, G. Carteni, and S. Cascinu, “Novel agents, combinations and sequences for the treatment of advanced renal cell carcinoma: when is the revolution coming?” Current Cancer Drug Targets, vol. 13, no. 3, pp. 313–325, 2013.
[23] B. I. Rini, G. Wilding, G. Hudes et al., “Phase II study of axitinib in sorafenib-refractory metastatic renal cell carcinoma,” Journal of Clinical Oncology, vol. 27, no. 27, pp. 4462–4468, 2009.
[24] O. Rixe, R. M. Bukowski, M. D. Michaelson et al., “Axitinib treatment in patients with cytokine-refractory metastatic renal-cell cancer: a phase II study,” Lancet Oncology, vol. 8, no. 11, pp. 975–984, 2007.
[25] B. I. Rini, B. Escudier, P. Tomczak et al., “Comparative effectiveness of axitinib versus sorafenib in advanced renal cell carcinoma (AXIS): a randomised phase 3 trial,” The Lancet, vol. 378, no. 9807, pp. 1931–1939, 2011.
[26] B. I. Rini, V. Grünwald, M. N. Fishman et al., “Axitinib for first-line metastatic renal cell carcinoma (mRCC): overall efficacy and pharmacokinetic analyses from a randomized phase II study,” Journal of Clinical Oncology. In press.
[27] P. Bhargava, B. Esteves, D. A. Nosov, O. N. Lipatov, A. A. Lyulko, A. A. Anischenko et al., "Updated activity and safety results of a phase II randomized discontinuation trial (RDT) of AV-951, a potent and selective VEGFR1, 2, and 3 kinase inhibitor, in patients with renal cell carcinoma (RCC)," *Journal of Clinical Oncology.* In press.

[28] R. J. Motzer, D. Nosov, T. Eisen, I. N. Bondarenko, V. Lesovoy, O. N. Lipatov et al., "Tivozanib versus sorafenib as initial targeted therapy for patients with advanced renal cell carcinoma: results from a phase III randomized," *Journal of Clinical Oncology.* In press.

[29] http://www.fda.gov/downloads/AdvisoryCommittees/CommitteesMeetingMaterials/Drugs/OncologicDrugsAdvisoryCommittee/UCM350075.pdf.

[30] E. Angervin, V. Grünwald, A. Ravaud, D. A. Castellano, C. C. Lin, J. E. Gschwend et al., "A phase I study of dovititinib (TK1258), an FGFR- and VEGFR-inhibitor, in patients with advanced or metastatic renal cell cancer (mRCC)," *Journal of Clinical Oncology.* In press.

[31] J. C. Welti, M. Gourlaouen, T. Powles et al., "Fibroblast growth factor 2 regulates endothelial cell sensitivity to sunitinib," *European Journal of Cancer,* vol. 30, no. 10, pp. 1183–1193, 2011.

[32] P. Mulders, R. Hawkins, P. Nathan et al., "Cediranib monotherapy in patients with advanced renal cell carcinoma: results of a randomised phase II study," *European Journal of Cancer,* vol. 48, no. 4, pp. 527–537, 2012.

[33] N. M. Tannir, Y.-N. Wong, C. K. Kollmannsberger et al., "Phase 2 trial of linifanib (ABT-869) in patients with advanced renal cell cancer after sunitinib failure," *European Journal of Cancer,* vol. 47, no. 18, pp. 2706–2714, 2011.

[34] M. Jermann, R. A. Stahel, M. Salzberg et al., "A phase II, randomised phase II study of AMG 386 in combination with sorafenib," *Clinical Cancer Research,* vol. 16, no. 14, pp. 3628–3638, 2010.

[35] D. Roulin, L. Waselle, A. Dormond-Meulew, M. Dufour, N. Demartines, and O. Dormond, "Targeting renal cell carcinoma with NVP-BEZ235, a dual PI3K/mTOR inhibitor, in combination with sorafenib," *Molecular Cancer,* vol. 10, article 90, 2011.

[36] B. Rini, C. Szczyluk, N. M. Tannir, P. Koralewski, P. Tomczak, A. Deptala et al., "AMG 386 in combination with sunitinib in patients with metastatic clear cell carcinoma of the kidney: a randomized, double-blind, placebo-controlled, phase 2 study," *Cancer,* vol. 118, pp. 6152–6161, 2012.

[37] M. B. Atkins, A. Ravaud, G. Gravis, K. Drosik, T. Demkow, P. Tomczak et al., "Safety and efficacy of AMG 386 in combination with sunitinib in patients with metastatic renal cell carcinoma (mRCC) in an open-label multicenter phase II study," *Journal of Clinical Oncology.* In press.

[38] V. Ramakrishnan, V. Bhaskar, D. A. Law et al., "Preclinical evaluation of an anti-α5β1 integrin antibody as a novel antiangiogenic agent," *Journal of Experimental Therapeutics and Oncology,* vol. 5, no. 4, pp. 273–286, 2006.

[39] S. Yazji, R. M. Bukowski, V. Kondagunta, and R. Figlin, "Final results from phase II study of volociximab, an α5β1 anti-integrin antibody, in refractory or relapsed metastatic clear cell renal cell carcinoma (mCCrCC)," *Journal of Clinical Oncology.* In press.

[40] C.-L. Law, K. A. Gordon, B. E. Toki et al., "Lymphocyte activation antigen CD70 expressed by renal cell carcinoma is a potential therapeutic target for anti-CD70 antibody-drug conjugates," *Cancer Research,* vol. 66, no. 4, pp. 2328–2337, 2006.

[41] E. Oosterwijk, D. J. Ruiter, and J. Hoedemaeker Ph., "Monoclonal antibody G 250 recognizes a determinant present in renal-cell carcinoma and absent from normal kidney," *International Journal of Cancer,* vol. 38, no. 4, pp. 489–494, 1986.

[42] Z. Liu, F. E. Smyth, C. Renner, E.-T. Lee, E. Oosterwijk, and A. M. Scott, "Anti-renal cell carcinoma chimeric antibody G250: cytokine enhancement of in vitro antibody-dependent cellular cytotoxicity," *Cancer Immunology, Immunotherapy,* vol. 51, no. 3, pp. 171–177, 2002.

[43] Y. Shao, Z. Gao, P. A. Marks, and X. Jiang, "Apoptotic and autophagic cell death induced by histone deacetylase inhibitors," *Proceedings of the National Academy of Sciences of the United States of America,* vol. 101, no. 52, pp. 18030–18035, 2004.

[44] P. Rocchi, R. Tonelli, C. Camerin et al., "p21Waf1/Cip1 is a common target induced by short-chain fatty acid HDAC inhibitors (valproic acid, tributyrin and sodium butyrate) in neuroblastoma cells," *Oncology Reports,* vol. 13, no. 6, pp. 1139–1144, 2005.
[54] H. Wang, W. Zhou, Z. Zheng et al., “The HDAC inhibitor depsipeptide transactivates the p53/p21 pathway by inducing DNA damage,” *DNA Repair*, vol. 11, no. 2, pp. 146–156, 2012.

[55] J. D. Hainsworth, J. R. Infante, D. R. Spigel, E. R. Arrowsmith, R. V. Boccia, and H. A. Burris, “A phase II trial of panobinostat, a histone deacetylase inhibitor, in the treatment of patients with refractory metastatic renal cell carcinoma,” *Cancer Investigation*, vol. 29, no. 7, pp. 451–455, 2011.

[56] G. Niu, K. L. Wright, M. Huang et al., “Constitutive Stat3 activity up-regulates VEGF expression and tumor angiogenesis,” *Onco-gene*, vol. 21, no. 13, pp. 2000–2008, 2002.

[57] A. Horiguchi, T. Asano, K. Kuroda et al., “STAT3 inhibitor WP1066 as a novel therapeutic agent for renal cell carcinoma,” *British Journal of Cancer*, vol. 102, no. 11, pp. 1592–1599, 2010.

[58] C. R. Ireson and L. R. Kelland, “Discovery and development of anticancer aptamers,” *Molecular Cancer Therapeutics*, vol. 5, no. 12, pp. 2957–2962, 2006.

[59] H. Ginisty, H. Sicard, B. Roger, and P. Bouvet, “Structure and functions of nucleolin,” *Journal of Cell Science*, vol. 112, no. 6, pp. 761–772, 1999.

[60] S. Soundararajan, W. Chen, E. K. Spicer, N. Courtenay-Luck, and D. J. Fernandes, “The nucleolin targeting aptamer AS1411 destabilizes Bcl-2 messenger RNA in human breast cancer cells,” *Cancer Research*, vol. 68, no. 7, pp. 2358–2365, 2008.