

Letter to the Editor

Experimental study on the prevention of liver cancer angiogenesis via miR-126. Promising results for targeted therapy

Dear Editor,

We read with great interest the paper written by Jing et al¹, recently published on Eur Rev Med Pharmacol Sci and titled "*Experimental study on the prevention of liver cancer angiogenesis via miR-126*", very interesting and promising topic based on miR-126 therapy.

Furthermore, recent issues reporting the exact identification of cellular signaling pathways playing a key role in the angiogenesis have led to development of new molecules for targeted therapies². This research is very important and could open future scenarios for the planning of targeted therapies coupling miR-126 and novel angiogenesis inhibitors³.

To date, due to novel drugs (Table I) and more pathological knowledge in the outcome of hepatocellular carcinoma (HCC), patients have improved also in so called "frail patients"⁴. Unfortunately, insufficient data are reported also in clinical trials in HCC patients who carrying concomitant HIV/HCV/HBV infections and liver metabolic disease⁵⁻⁸.

Considering the heterogeneous aspects of HCC and to enhance therapeutic efficacy, overcome drug resistance and reduce toxicity, combination of antiangiogenic drugs with antineoplastic chemotherapy (AC), radiotherapy or other targeted drugs as miRNAs, have been evaluated⁹. To date, there is no designed approach to determine which patients are most likely responsive to a given type of antiangiogenic treatment based on miRNAs molecules.

However, another aspect will be focusing on the miRNAs and in particular miR-126 is the efficacy of their detection¹⁰. Moreover, if the detection of miR-126 is routinely incorporating into clinical practice, knowledge concerning the predictive value of this test will eventually enable individual therapy in HCC patients¹¹.

Several approaches to consider the quality of the treatments for suitable cost-effective models into healthcare systems are ongoing consideration in HCC managements¹².

Finally, the results obtained by Jing et al¹ are interesting, but, in our opinion, need of more confirmation studies, considering the high incidence of HCC and its tissue heterogeneity in human model.

In addition, we think that the recent progress in genetics have provided exceptional opportunities to identify prognostic and predictive markers of efficacy of antiangiogenic treatments¹³. Genetic markers could be a tool to identify patients who will benefit from this targeted therapy, and exclude patients with high risk to develop severe toxicity¹⁴.

We know that this kind of translational studies are strongly recommended to have more and more information on HCC with the aim to improve the efficacy of treatment and quality of patients life¹⁵. This field of research might be able to accelerate the translational research into the routine therapies.

Conflict of interest

The Authors declare that they have no conflict of interests.

Table I. Most common antiangiogenic molecules used in HCC.

Drugs (Alias)	Typology	Targets	HCC therapeutic indication and study phase
<i>Bevacizumab</i>	Monoclonal antibody	VEGF	Phase II study in combination with Erlotinib (NCT00881751)
<i>Brivanib</i>	Small molecule Multikinase inhibitor	VEGFR, FGF	Advanced stage, Phase III study in patients who failed Sorafenib (NCT00825955), and first line (NCT00858871)
<i>Cabozatinib</i>	Small molecule Multikinase inhibitor	c-MET, RET, VEGFR1-3, c-KIT	Preclinical study, yet.
<i>Cediranib</i>	Small molecule Multikinase inhibitor	VEGFR 2, PDGFR, c-Kit	Advanced stage, phase II study
<i>Everolimus</i>	Small molecule	m-TOR	Not registered; phase III study did not show significant efficacy
<i>Lenvatinib</i>	Small molecule Multikinase inhibitor	VEGFRs 1, 2 and 3, FGFRs 1 PDGFR α , RET, KIT	Not registered; phase III trial underway (NCT01761266)
<i>Linfantib</i>	Small molecule Multikinase inhibitor	VEGFR 2, PDGFR families	Advanced stage, phase III study first-line vs sorafenib (NCT01009593)
<i>Nintedanib</i>	Small molecule Multikinase inhibitor	VEGFRs 1, 2 and 3, FGFRs 1 PDGFR	Phase design study vs sorafenib (NCT00987935 and NCT01004003)
<i>Ramucirumab</i>	Monoclonal antibody	Selective VEGFR 2	Not registered; phase III study with conflicting results
<i>Refametinib</i>	Small molecule Mitogen-activated protein kinase inhibitor	MEK 1-2	Advanced stage, phase II study in patients who failed sorafenib (NCT01915589) and in combination with sorafenib (NCT01915602)
<i>Regorafenib</i>	Small molecule Multikinase inhibitor	VEGFR1-3, c-KIT, TKI g -like, EGF-like 2, PDGF 2, FGF 1, RET, RAF-1, BRAF, MAPK	Not registered; phase III study underway recruiting patients who progressed under sorafenib (NCT01774344).
<i>Sorafenib</i>	Small molecule Multikinase inhibitor	VEGFR 2, PDGFR, c-Kit, BRAF	Advanced stage; not recommended for use in adjuvant treatment
<i>Trebananib</i>	Small molecule Angiogenesis inhibitor	Tie-2 (Angiopoietin)	Not registered; phase II study no showed improvement of OS.
<i>Vatalanib</i>	Small molecule Angiogenesis inhibitor	VEGFRs 1, 2 and 3, PDGFR c-FMS	Advanced stage, phase II in combination with Doxorubicin.

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