

Editorial – HCC in HCV patients and the direct acting antivirals: is there really a link?

A. MARRONE¹, G. FRANCI², A. PERRELLA³, R. NEVOLA¹, A. CHIANESE²,
L.E. ADINOLFI¹, F.C. SASSO¹, L. RINALDI¹

¹Department of Advanced Surgical and Medical Sciences, University of Campania Luigi Vanvitelli, Naples, Italy

²Department of Experimental Medicine, University of Campania Luigi Vanvitelli, Naples, Italy

³Immunological and Neurological Infectious Diseases Unit, Azienda Ospedaliera dei Colli, Cotugno Hospital, Naples, Italy

The onset of hepatocellular carcinoma (HCC) in hepatitis C virus (HCV) patients who achieved the sustained virological response (SVR) after treatment with direct acting antivirals (DAAs) is still a matter of study and debate.

Currently, most of the literature is still focusing attention on the assessment of epidemiological data as well as the incidence of HCC during follow-up, compared to the annual mean rate in cirrhotic patients¹⁻⁹. Globally, among the results of several studies^{3,4,10-13}, a trend towards a progressive reduction or stability of HCC risk with SVR seems to be prevalent. Furthermore, scholars¹⁴⁻¹⁷ have shown, especially in the early stages after treatment, the appearance of unexpectedly large HCC lesions. More recently, some preliminary papers are showing a late HCC *de novo* occurrence in patients after more than 18 months of SVR, suggesting possible other mechanisms subtended to HCC occurrence even in absence of HCV chronic inflammation¹⁸. Therefore, these contrasting epidemiological data clearly suggest that mechanisms in carcinogenesis are still unclear, requiring further investigations. Several papers have identified some variables most frequently associated with the appearance of HCC *de novo*, such as the value of platelets, liver stiffness, portal hypertension, alpha-feto protein value, alcohol consumption, metabolic factors or the most advanced stage of cirrhosis¹⁹⁻²⁵. However, these elements appear to be major risk predictors but are not attributable to direct carcinogenic effect. The possible indirect implication of DAAs was immediately discussed.

Our group, in a recent prospective multicenter work²⁶, has highlighted a predominance of HCC *de novo* in patients who have been treated with sofosbuvir without ribavirin in the 2015-2017 period. Our hypothesis is based on the lack of immunomodulatory protective effect of ribavirin²⁷ and on the possible damage on the mitochondrial RNA of sofosbuvir²⁸.

Likewise, some authors focused on the reduced immunosurveillance induced by early and rapid viral clearance, which may partly explain the controversial relationship between eradication and HCC occurrence. Researchers^{29,30} have highlighted the possible role of some cytokines as carcinogenesis co-factors. To date, there is no direct evidence of an involvement of DAAs in the carcinogenesis.

However, it should also be considered that we still do not have details on the new antivirals according to their possible immunomodulatory activity. Indeed, in order to infection disease, the antivirals are well known to exert some immunomodulatory activity. The most interesting immunomodulating activity is for two sub-set of antivirals as the protease inhibitors as, for instance, fosamprenavir or darunavir^{31,32} and maraviroc³³ being involved in the modulation of the angiogenesis, tumor growth and invasion, inflammation, antigen processing and presentation, cell survival. Instead, concerning HCV antivirals, ribavirin has been demonstrated to exert a kind of immunomodulatory activity markedly reducing macrophage activation and diminishing Th2 cytokine production while preserving Th1 cytokine production and stimulating humoral immune response^{34,35}. All these studies and the current literature debate about HCC *de novo* onset after SVR following new DAAs therapy clearly suggest that most of the drugs, even if to be considered safe, may exert some important effect on host that should be considered.

Despite the fact that the DAA therapies are able to remove the HCV from the patients, the epigenetic impact on cells persist also in absence of the virus itself. It is becoming emergent the relationship and

connection among these alterations and the risk of development of HCC during the lifetime³⁶. The HCV infection determines an epigenetic furrow with a determinate signature. This signature is made of DNA methylation and histone modifications. Hamdane et al³⁷ highlights how the acetylation of lysine 27 on the histone 3 (H3K27Ac), due to the HCV, promote the alteration of gene expression involved in the Protein Kinase B (AKT), mammalian Target of rapamycin (mTor), Tumor Necrosis Factor alpha (TNF- α) pathways with an increase of HCC risk. Indeed, the overexpression of the SPHK1 gene (mediator of sphingolysin phosphorylation) and SP1 (regulator of inhibition of apoptosis and promotion of proliferation) has been demonstrated to play a crucial role in the TNF- α signalling pathway³⁷. HCV infection reflects on DNA methylation alteration. As double side of the same coin, DNA methylation is responsible in gene activation and silencing. HCV infection can determine in the same time hyper and lower-methylated region. Zheng et al³⁸ highlights how the decrease in specific region, such as the interspersed nuclear element-1 (LINE-1) is a direct consequence of HCV infection. This aspect is due to natural kill cell-dependent innate immune response activation with an aberrant DNA methylation in hepatocytes³⁹. All together the information opens the scenario of epigenetic relationship among DAA treated patients and HCC risk.

An additional pathogenetic mechanism can be determined by gut microbiota which has been demonstrated to have an important connection in the inflammation, liver fibrosis, and HCC⁴⁰. Indeed, a recent study showed that HCV eradication with DAAs in patients with cirrhosis is associated with a modification of the gut microbiota, which correlates with fibrosis and inflammation, but does not improve intestinal barrier function⁴¹.

HCC is still today a high mortality oncological illness⁴² with multifactorial aspects related to genetics^{43,44} to virology⁴⁵ and metabolic factors^{46,47}.

The research for effective medical therapies is subject to an understanding of these mechanisms. The possible indirect role of DAAs in carcinogenesis may represent a further food for thought.

Conflict of Interest

The Authors declare that they have no conflict of interests.

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