

# **New Therapies, Markers and Therapeutic Targets in HCV Chronic Infection, and HCV Extrahepatic Manifestations.**

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## **Abstract**

**More than 180 millions of subjects in the world are infected by Hepatitis C Virus (HCV), and about 20% of them with HCV chronic infection progress to cirrhosis.** Furthermore, numerous HCV extrahepatic manifestations have been reported in up to 74% of patients, as mixed cryoglobulinemia, lymphomas, rheumatic disorders, autoimmune thyroiditis, hypothyroidism, papillary thyroid cancer, and type 2 diabetes. Advances in understanding the HCV life cycle, and the inflammatory processes (involving a complex network of cytokines and chemokines) associated with HCV chronic infection, have led to substantial advancements in therapy. **The combination of ribavirin and PEGylated interferon- $\alpha$  was the standard of therapy for HCV chronically infected patients in the last decades.**

**However,** interferon has limited effectiveness and is associated with severe adverse effects. **Recently, direct-acting antivirals (DAAs) that act as inhibitors of N5SA, or polymerase, or protease have been shown to result in shorter duration of therapy, better efficacy and tolerance, with respect to ribavirin and PEGylated interferon- $\alpha$ .** Circulating CXCL10 levels, and **the interleukin(IL)-28B gene polymorphisms,** are associated with the **success of the therapy both with DAAs or ribavirin and PEGylated interferon-alpha. New DAAs targeting the HCV at various molecular levels have been developed to eradicate HCV.** Moving to **interferon-free** therapies should offer new treatments for resistant HCV genotypes, and for ineligible patients or patients failing to respond to prior therapies.

**Many efforts have been made to understand the factors that are involved with clearance of HCV to personalize the therapy for each patient, with the aim to reduce side effects, increasing the sustained virologic response rate, and to prevent the progression of the disease.**

## EDITORIAL

**More than 180 millions of subjects in the world are infected by Hepatitis C Virus (HCV), and about 20% of them with HCV chronic infection progress to cirrhosis [1].**

Furthermore, numerous extrahepatic manifestations (HCV-EHMs) have been reported in up to 74% of patients [2-5]. HCV strategies that limit or delay the initiation of innate antiviral responses are important in determining the final outcome of the infection [6]. High rate of replication and error prone genome replication machinery enable HCV to evade immune recognition. This delay of an effective immune response allows HCV to establish widespread infection in liver and extra-hepatic tissues and cells. Changing viral epitopes induces a failure of the adaptive immune response. Moreover, neutralizing antibody epitopes may be hidden by decoy structures, lipoproteins and glycans [7, 8]. Also, T cell responses fail due to changing viral epitope sequences; while the phenomenon of exhaustion is probably evolved to limit immune-mediated pathology [9]. As a result, immune-mediated clearance of HCV infection is occurring only in about 20% of people, by innate and adaptive immune mechanisms [10].

**A complex orchestration of cytokines and chemokines coordinate immune response, both innate and adaptive, in the initial phases of HCV infection,** and plays a pivotal role in controlling viral replication and liver damage [11]. Among cytokines, it is well known that the production of and/or response to interferons (IFNs) are very important in the induction of the immune response during chronic HCV infection (CHC) [12, 13], and several studies have shown that patients with CHC have a deregulated IFNs response [14]. In fact, patients who already have a high level of endogenous IFN-stimulated genes (ISGs) expression do not achieve viral clearance or have a poor response to treatment with IFN- $\alpha$  [15]. The viral and/or host factors that are responsible for the greater baseline ISGs expression in certain HCV patients remain to be determined [16]. Recently, it has been shown that type III IFNs, and in particular the new discovered IFN- $\lambda 4$ , play a dominant role in driving ISGs response and in contributing to the viral clearance or persistence [17-19].

A predominance of the Th1 immune response (and related cytokines/chemokines) has been shown in CHC and in HCV-EHMs [20, 21]. **Interferons, and chemokines inducible by interferon- $\gamma$  [(C-X-C motif) ligand (CXCL)9, -10 and -11] recruit inflammatory cells into the hepatic parenchyma (when the infection is not controlled) producing chronic inflammation and fibrosis of the liver, that may results in hepatic cirrhosis [22-25]. A complex dysregulation of cytokines and chemokines response is associated with HCV systemic manifestations [mixed cryoglobulinemia (MC), lymphomas, autoimmune diseases of the thyroid (AITD), type 2 diabetes], and involves overall the Th1 cytokines/chemokines [23, 26]. HCV escapes immunity interfering at various levels with cytokines/chemokines, and inducing a Th2/Tc2 immune response [26, 27]. The administration of interferon- $\alpha$  can induce the HCV clearance during CHC, and can revert the progression to cirrhosis, reducing CXCL10 levels [25, 28]. It has been suggested that agents directed to neutralize CXCL10 could increase the responsiveness of patients to traditional therapies of HCV infection, simultaneously reducing inflammatory immune cell activation [29]. Many studies have shown that both circulating CXCL10, and polymorphisms of interleukin (IL)-28B, could be used as prognostic markers of efficacy of HCV therapies [30-33].** Other studies have also shown that HCV clearance by direct-acting antiviral (DAAs) therapies decreases circulating CXCL10 levels [34].

HCV infection regulates a number of microRNAs (miRNAs), which are able to exert an effect on liver biology and pathology [35, 36]. In HCV infected hepatocytes miRNAs can directly regulate HCV replication through interaction with the HCV genome [37]. Moreover miRNAs induced by HCV can indirectly control critical virus-associated host pathways, inducing liver fibrosis, cirrhosis, and/or hepatocellular carcinoma [38, 39]. Recently, circulating miRNAs are emerging as biomarkers for HCV associated disorders. Considerable efforts have been employed to investigate the change in the circulating miRNA pattern **in HCV infection and related diseases** [40, 41]. **Distinctive circulating miRNA patterns are associated with**

**HCV infection and HCV-related hepatic diseases [42], suggesting that miRNAs are non-invasive biomarkers to evaluate the diagnosis and prognosis of these disorders [43].**

Treatment for HCV infection has recently progressed from poorly tolerated IFN- $\alpha$  therapy and with very low cure rates, to highly effective oral DAAs with cure rates above 90% for almost all patients, and with little adverse effects [44-46]. Understanding of the viral lifecycle, with recognition of targets that could be inhibited by small molecules, **has permitted the production of DAAs, that inhibit protease, non-structural 5a (NS5A), and nucleotide and non-nucleotide polymerase [47-49].** Initially DAAs have been used **with Pegylated-Interferon- $\alpha$  (Peg- IFN) and Ribavirin (RBV),** and subsequently **in** combination without the need for IFNs [50]. Rational DAAs combinations have overcome the major challenge of rapid emergence of drug resistance [51]. Second-generation DAAs agents in each class have further improved safety and efficacy profiles, reducing drug-drug interactions and adverse side effects [52-54].

In the new era of DAAs therapy in which elimination of HCV infection is a real possibility, HCV infected cell culture models are important for the identification of therapeutic targets, testing candidate drugs, and profiling of therapeutic strategies [55-58]. The development of protocols to grow HCV in culture and generate hepatocyte cell lines from specific individuals holds great promise to investigate the mechanisms exploited by the virus to spread the infection and the host factors critical for HCV replication and propagation, or resistance to infection [59-63]. These models could be used for the development of drugs targeting host factors essential for virus replication, holding great promises in further increasing treatment efficacy [64].

The most common HCV-EHMs of HCV infection is MC syndrome (MCs), an immune-complex **mediated vasculitis, involving joints, skin, peripheral nerves, and internal organs.** In **the** majority of individuals, MCs shows a mild, slow-progressive clinical course needing only symptomatic treatments [65-68]. However the etiologic therapy was considered the first-line option in MCs patients and, in the past two decades, antiviral treatment with **RBV plus Peg-IFN represented the standard of care** [69]. Rapidly progressive, diffuse MC

vasculitis with multiple organ involvement may be successfully treated with aggressive immunosuppressive and anti-inflammatory therapies, mainly based on cyclophosphamide or rituximab, high dose corticosteroids, and plasmapheresis [70]. The recent introduction of DAAs substantially changed the treatment of HCV infection. DAAs anti-HCV therapy seem to be safe and effective in patients with MC and MCs from the virological and clinical points of view, thus confirming the key role played by HCV eradication in inducing MC remission [71-73]. **The combination, or a sequence, of antiviral and immunosuppressive treatments has been shown to be useful in the therapy of MCs patients with major clinical manifestations. In the clinical practice, treatment of MCs needs to be personalized for each patient in relation to clinical symptoms and the severity of the diseases [74-76].**

On the bases of epidemiological data, biological investigation, as well as clinical observations, HCV is increasingly recognized to be involved in the pathogenesis of a variety of histological **types of B-cell non-Hodgkin' s lymphoma (B-NHL) [77-80]. To understand the mechanisms related to HCV persistence, and lymphomagenesis, is necessary to develop new therapies with the aim to prevent and to treat B-NHLs [81, 82].** It has been also observed that at least some types of HCV-related indolent B-NHLs disappear after successful antiviral treatment, strongly reinforcing the hypothesis of a HCV-induced lymphomagenesis [83]. The role of DAAs in the prevention and treatment of HCV associated lymphoproliferation remains to be clarified [84, 85].

Other HCV-EHMs are various rheumatic disorders (mainly arthritis, sicca syndrome, osteosclerosis, etc) [86-91]. The management of HCV-associated rheumatic diseases is particularly difficult **because of the coexistence of complex immunological disorders and HCV infection, and needs to be personalized in each patient [92-94].**

The occurrence of autoimmune thyroiditis, hypothyroidism, papillary thyroid cancer and type 2 diabetes is also more frequent in CHC: these disorders can significantly impact the quality of

life of HCV patients, the course of the disease, and the effect of therapies [95-98]. The role of DAAs in the prevention and treatment of these endocrine disorders remains to be clarified.

In this issue, we focus on the recent advancements in the viral and host factors influencing their interplay, highlighting current knowledge in the inflammatory processes (and the inherent network of cytokines and chemokines) associated with CHC, and illustrate the multifaceted HCV-EHMs, that heavily affect the quality of life of HCV patients. We then describe how the development of reliable systems to propagate the virus in vitro and the identification of small molecules targeting key steps of HCV replication have led to substantial advancements in the therapy. Finally, we give an overview of current approaches for treatment and of novel DAAs targeting **various stages of the life of HCV** and holding promise **to eradicate HCV infection**. Moving to IFN-free therapies should offer new treatments for resistant HCV genotypes, and for ineligible patients or patients failing to respond to prior therapies. **The knowledge of viral and host factors associated with the clearance of HCV is very important to personalize the therapy in each patient, increasing the sustained virologic response rates, preventing the progression of hepatic disease, and reducing adverse side effects.**

**Conflict of Interest**

The authors have no conflict of interests to declare.



## **Acknowledgements**

We thank the contributing authors, co-authors, and reviewers for sharing their valuable time and expertise to put this issue together.

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