



2016 Hepatitis C Virus: Global view

Analysis of peripheral blood dendritic cells as a non-invasive tool in the follow-up of patients with chronic hepatitis C

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Abstract

Hepatitis C virus (HCV) has a high propensity to establish chronic infections. Failure of HCV-infected individuals to activate effective antiviral immune responses is at least in part related to HCV-induced impairment of dendritic cells (DCs) that play a central role in activating T cell responses. Although the impact of HCV on DC phenotype and function is likely to be more prominent in the liver, major HCV-induced alterations are detectable in peripheral blood DCs (pbDCs) that represent the most accessible source of DCs. These alterations include numerical reduction, impaired production of inflammatory cytokines and increased production of immunosuppressive IL10. These changes in DCs are relevant to our understanding the immune mechanisms underlying the propensity of HCV to establish persistent infection. Importantly, the non-invasive accessibility of pbDCs renders the analysis of these cells a convenient procedure that can be serially repeated in patient follow-up. Accordingly, the study of pbDCs in HCV-infected patients during conventional treatment with pegylated interferon and ribavirin indicated that restoration of normal plasmacytoid DC count may represent an additional mechanism contributing to the efficacy of the dual therapy. It also identified the pre-treatment levels of plasmacytoid DCs and IL10 as putative predictors of response to therapy. Treatment of chronic HCV infection is changing, as new generation direct-acting antiviral agents will soon be available for use in interferon-free therapeutic strategies. The phenotypic and functional analysis of pbDCs in this novel therapeutic setting will provide a valuable tool for investigating mechanisms underlying treatment efficacy and for identifying predictors of treatment response.

Key words: Hepatitis C virus; Peripheral blood dendritic cells; Cytokines; Peg-interferon; Ribavirin

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Core tip: Dendritic cells (DCs) are professional antigen-presenting cells that play a primary role in the activation and coordination of primary immune responses. In this review we will illustrate and discuss the emerging understanding of DC impairment occurring in patients with chronic hepatitis C virus (HCV) infection and the impact of therapy on DCs. Particular attention will be paid to HCV-induced alterations of DCs in the peripheral blood, as the non-invasive accessibility of these cells renders their analysis a convenient procedure that can be serially repeated in patient follow-up.

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INTRODUCTION

One of the main features of the hepatitis C virus (HCV) is its high propensity to establish chronic infections that is related to impaired function and a decreased number of HCV-specific T cells^[1,2]. Given the important role of dendritic cells (DCs) in activating T cell responses^[3], DCs have become the target of investigation demonstrating that failure of HCV-infected individuals to activate effective antiviral immune responses is at least in part the result of HCV-induced impairment of DC function. Because of the difficulties in gaining access to liver DCs, although the liver is the primary site for HCV replication most studies addressing DC involvement in chronic HCV infection have focused on peripheral blood DCs (pbDCs). These studies provided useful information on the impact of HCV on DCs, the contribution of DCs to the pathogenesis of hepatitis C, the effects of therapy on DCs and DC features predictive of response to treatment.

CHRONIC INFECTION WITH HEPATITIS C VIRUS

Epidemiological aspects and natural history

HCV infection has a worldwide distribution and has been recognized as a major cause of chronic hepatitis, cirrhosis and end-stage liver disease. According to the 2013 World Health Organization report, about 150 million people are chronically infected by HCV worldwide and more than 350000 people die every year due to HCV-related complications^[1].

Up to the 1990's, the principal routes of HCV infection, were blood transfusion, unsafe injection procedures and intravenous drug use. Taken together these routes are estimated to be responsible for approximately 70% of chronic cases in developed countries. Currently screening of blood products for HCV by means of enzyme immunoassays (EIA) and nucleic acid testing has virtually eradicated transfusion-associated hepatitis C. Similarly, in the developed world, new HCV infections are infrequently related to unsafe medical or surgical procedures and spread among the community of people using intravenous drugs now accounts for the vast majority of incident cases. Other invasive behaviors, such as tattooing and acupuncture with unsafe materials are also implicated in occasional HCV transmissions. The risk of perinatal and of heterosexual transmission of HCV is low while male homosexual activity has become an important transmission route in Western Countries^[4]. Conversely, the situation is quite different in resource-poor countries, where lack of public awareness and continuous use of unsafe medical tools still account for a considerable proportion of new HCV infections.

HCV is a positive strand RNA virus characterized by high sequence heterogeneity. Seven HCV genotypes, numbered 1 to 7, and a large number of subtypes have been described^[5]. Genotypes and subtypes (which are identified by lowercase letters), differ among themselves by about 30% and 20% of their sequences respectively.

Acute hepatitis C is rarely severe, and symptoms occur in 10% to 50% of cases^[4]. The incidence of acute HCV infection has decreased and it is about 1/100000 per year but this figure is probably underestimated because it mainly refers to symptomatic patients. Progression to chronic infection occurs in about 75% of cases but again, the estimate is very roughs mainly due to difficulties in the diagnosis of the acute infection^[6] and possibly to difference related to the route of infection. For example, in a small outbreak of acute hepatitis C (genotype 2c) among subjects who took part in pharmacokinetics studies, chronicity occurred in less than 50% of subjects and some cases of very late clearance (up to 24 mo from the date of infection) were also described. In this latter case, unsafe procedures of blood sampling were supposed to be responsible for HCV infection^[7].

Chronic hepatitis C is characterized by a variable degree of inflammation and with variable rates of fibrosis progression. Only exceptionally, HCV infection clear spontaneously in the chronic stage. Depending on the viral genotype, between 15% and 45% of infected cases will spontaneously resolve the infection within six months of infection without any treatment whilst the remaining proportion of cases will maintain the virus for years, with a risk of developing liver cirrhosis within 20 years up to 15%-30%^[1]. On average, 20%-30% of patients develop cirrhosis over 20-30 years of infection^[6]. The risk of developing

hepatocellular carcinoma (HCC) in patients with HCV liver cirrhosis is approximately 1% to 5% per year. Patients with a diagnosis of HCC have a 33% probability of death during the first year of diagnosis^[8]. Overall the standardized mortality rate in anti-HCV-positive persons ranges from 1.6 to 4.5 and was as high as 25 in a recent study from Scotland^[9].

Hepatitis C progression to cirrhosis is highly variable depending on the presence of cofactors capable of accelerating the fibrotic process. Proven cofactors for fibrosis progression include older age at infection, male gender, chronic alcohol consumption, obesity, insulin resistance and type 2 diabetes, and immunosuppression (such as that occurring after solid organ transplantation and in untreated HIV infection)^[10]. The clinical management of these cofactors represents a mainstay of HCV treatment, especially because many of these cofactors are associated with poor response to clinical treatment.

Clinical management

The primary goal of HCV therapy is to cure the infection, which is generally associated with the resolution of liver disease in patients with less advanced disease, that is, in patients without cirrhosis. However, patients with cirrhosis who have eradicated viral infection remain at risk of severe complications, mainly the development of HCC^[11] and therefore need to continue active ultrasonographic follow up for early detection of HCC.

The infection is cured in more than 99% of patients who achieved a sustained virological response (SVR), defined as undetectable HCV RNA 24 wk after treatment completion^[9]. Until recently, the only therapeutic option for chronic HCV infection was the combination of pegylated interferon (pegIFN) plus ribavirin (RBV), also known as dual therapy. With this regimen, patients infected with genotype 1 had SVR rates ranging from 40% in North America to 50% in Western Europe. Genotype 4 too, is characterized by low SVR rate. Higher SVR rates were achieved in patients infected with other HCV genotypes. For genotype 2, SVR rates accounted for about 80% while for genotypes 3, 5 and 6 the success rate appears to be lower. Both pegylated IFN molecules, pegylated IFN α 2b (1.5 mcg/kg/wk) and pegylated IFN α 2a (180 mcg/wk) can be used. Ribavirin should be given at a weight based daily dose (15 mg/kg) for genotypes 1, 4, 5 and 6 whereas, for genotype 2 and 3, a flat dose of 800 mg/d is indicated^[9].

Although they do not target a specific HCV protein or nucleic acid, both pegIFN and RBV exert their antiviral activity through induction of various antiviral proteins and through different immunomodulating actions. Dual therapy has been largely used for all genotypes but it is characterized by both limited efficacy and poor tolerability^[9,12] and these limitations have stimulated research to explore the feasibility of new therapeutic tools. HCV interferon-dependent

therapies rely on host factors such as IL28B polymorphism, liver fibrosis stage, and prior peg/IFN-RBV history to predict treatment response^[13,14].

Recently, improvement in our knowledge of the molecular biology of the HCV replication life cycle led to the discovery of several molecules that specifically block various viral proteins^[15,16]. These compounds are called direct-acting antiviral agents (DAA) and target different nonstructural proteins involved in the HCV life cycle, including the NS3/4A protease, the NS5B polymerase and the NS5A protein. There are 4 classes of DAAs: NS3/4A protease inhibitors; NS5B nucleos(t)ide inhibitors; NS5B nonnucleos(t)ide inhibitors; NS5A inhibitors^[17]. The first DAAs approved for the treatment of chronic HCV infection were the protease inhibitors (PI) telaprevir and boceprevir. Efficacy of these two first PI generation is limited to genotype 1, being genotype 1a less responsive than genotype 1b. The addition of PI to dual therapy significantly improved the efficacy of treatment for genotype 1 HCV infection but at the expenses of worsened side effect profile with the potential for severe adverse events, mainly hematological complications^[18-21].

All naive patients with compensated disease due to HCV should be considered for therapy. Treatment should not be deferred in case of significant fibrosis (METAVIR score F3-F4). For genotype 1 the addition of telaprevir or boceprevir to the dual therapy is the approved standard of care. Duration of therapy ranges between 24 and 48 wk depending on the rapidity of viral clearance. However, these triple-therapy regimens are complex, with variable viral monitoring and stopping rules. All patients with cirrhosis should be treated for 48 wk. Only selected patients with contraindication to boceprevir or telaprevir and/or with highly likelihood of SVR should be administered dual therapy^[9].

The combination of pegIFN and RBV is the approved standard of care for chronic hepatitis C, genotypes 2, 3, 4, 5 and 6. Dual-treatment duration should be tailored to on-treatment virological response. The likelihood of SVR is directly proportional to the speed of HCV RNA disappearance. Treatment should be stopped at week 12 if the HCV RNA decrease is less than 2 log UI/mL and at week 24 if HCV RNA is still detectable^[9]. In patients with rapid virological response (undetectable HCV RNA at week 4), duration of treatment should be 24 wk but, for genotype 1 and 4, 24 wk of therapy may be sufficient only in the presence of low viral load (< 400000 UI/mL) at baseline; the remaining patients should be treated for 48 wk. Patients with later virological response should be treated for 48 wk, as well^[9].

The protease inhibitor, Simeprevir and the nucleos(t)ide polymerase inhibitor, Sofosbuvir have been the first DAAs approved for HCV treatment. They were shown to be safer and more effective compounds compared with interferon-based therapy. Subsequently

several other DAAs have also been approved and many other compounds from all DAA families are still under evaluation. Therefore, interferon-free strategies for the therapy of HCV chronic infection will be available in the near future with many advantages in terms of tolerability. This is the reason why many doctors and patients are now choosing to defer treatment rather than to proceed with dual or triple therapy.

IMMUNE EFFECTOR MECHANISMS IN THE CONTROL OF HCV INFECTION

The role of T cells

The mechanisms causing the high propensity of HCV to establish chronic infections are not clearly defined, but it is believed that alterations of the antiviral immunity play a central role in determining the clinical outcome of infection. Indeed, the presence of strong and multi-specific T cell mediated antiviral responses is a hallmark of positive outcome of infection. It has been previously shown that a higher proportion of subjects who spontaneously cleared the HCV infection in the past have T cell proliferative responses to HCV peptides in comparison with patients with established chronic infection^[22,23]. Furthermore, the magnitude of these responses was also greater in past responders^[22]. In addition to different antigen-specific T cell proliferation profiles, different clinical outcomes of the HCV infections also correlate with different profiles of antigen-specific T cell cytokine production. T cells producing high amounts of IFN γ in response to multiple HCV antigens can be detected (1) in subjects with acute HCV infections as they progress to spontaneously clear the virus but not in subjects with acute HCV infection who become chronic^[24]; (2) in subjects with past resolved HCV infection but not in patients with established chronic infection^[23]; (3) in subjects with chronic HCV infection who respond to IFN α plus Ribavirin and pharmacologically clear the virus but not in those who do not respond^[25]; and also (4) in chronic HCV treatment responders with faster response kinetics (fast responders) but only to a lesser extent in slow responders^[26]. Furthermore, the presence of virus-specific T cell mediated responses can be observed even in the absence of anti-HCV antibodies in subjects who cleared the infection years before^[23,27] and also in HCV-negative children born from HCV-positive mothers^[28], where detectable HCV-specific T cells show increased production of IFN γ and reduced production of IL10 in response to HCV peptides in comparison to T cells obtained from the infected mothers, revealing a potential use of antigen-specific T cell detection as a powerful diagnostic marker to identify cases of viral exposure even in the absence of antibodies detectable with currently available methods.

The role of IFN γ and other cytokines

IFN γ is the most powerful T cell-derived antiviral

factor and inducer of pro-inflammatory responses (reviewed in^[29,30]). Amongst its functions, IFN γ boosts the cytotoxic activities of CD8⁺ T cells and NK cells, favours the differentiation of Th1 cells, enhances the killing activities mediated by phagocytes of the innate immunity, upregulates the expression of both class I and class II MHC molecules on by-stander cells and also stimulates activated B cells to perform antibody isotype switch towards IgG classes that are favourable for opsonisation and phagocytosis, complement activation and antibody-dependent cell cytotoxicity. Furthermore, besides its immune activities IFN γ can-like the other interferons, alpha, beta and lambda-act directly on infected cells, inducing intracellular genetic programs of direct antiviral response involving the expression of several interferon-stimulated genes (ISGs) that directly target viral metabolites suppressing viral replication. Infected cells can therefore be purged of the virus without undergoing immune-mediated killing, and by-stander cells can achieve an antiviral state that renders them refractory to infection. The production of IFN γ is facilitated by innate pro-inflammatory cytokines such as IL12, IL15 and IL18, which are mainly produced by activated antigen-presenting cells.

IL10 and other inhibitory pathways

If the production of IFN γ develops hand-in-hand with and favours the control of the infection, the production of anti-inflammatory IL10 follows different kinetics. In contrast with the pro-inflammatory and antiviral role of IFN γ , IL10 is one of the strongest endogenous immune suppressing agents physiologically produced by multiple cell subsets of both the innate and the adaptive immune system, including antigen-presenting cells (monocytes, dendritic cells) and subsets of antigen-specific T cells (reviewed in^[31,32]). The physiological role of IL10 is to shut down no longer required immune responses, thus preventing continuous immune activation and immune-mediated tissue damage. Functionally speaking, IL10 contrasts several functions exerted by IFN γ , effectively suppressing antigen processing and presentation by antigen-presenting cells and T-cell-mediated antiviral functions, both direct and indirect. IL10 can suppress the production of almost all pro-inflammatory cytokines and chemokines^[32], and can direct the differentiation of anergic T cells from naïve precursors, thus contributing to the development of impaired T-cell responses^[33-35].

The production of IL10 can be induced in combination with other immunological negative check points that share functional similarities, such as the Programmed-Death-1 (PD1) pathway. Several groups have shown that in patients with established chronic HCV infection HCV-specific T cells manifest with an immune exhausted phenotype^[36-38] characterized by suppressed antigen-specific proliferation and cytokine production together with increased expression of PD1

and other T-cell activation markers that act as inhibitory immune check points, such as T-cell immunoglobulin mucin-3 (TIM3), lymphocyte activation gene-3 (LAG3), cytotoxic T-lymphocyte-associated protein 4 (CTLA4) and CD244^[39]. Furthermore, antigen-specific T-cell exhaustion is not specific of chronic HCV infections only, but it can be observed during other chronic viral infections, including HBV^[40-44] and HIV^[45-47]. Monocytes expressing PD1 were shown to produce IL10 in response to PD1 engagement during HIV infections^[48], with the consequence of suppressing HIV-specific immunity. Furthermore, in alcoholic liver disease the production of IL10 was correlated with reduced expression of IFN γ and increased expression of the immune inhibitors PD1 and TIM3 on several immune subsets in response to bacterial stimulation^[49]. Several dysfunctions associated with immune exhaustion can be successfully reversed using blocking antibodies targeting these hyper-expressed immune inhibitory receptors and their respective ligands^[45-47,50-52], and immunotherapeutic protocols using these antibodies are currently being evaluated in clinical studies for a plethora of medical conditions, including viral infections and cancer^[53,54].

In the context of acute HCV infections, strong HCV-specific IL10 responses develop only during later stages of the acute infection in subjects who develop chronicity^[24], possibly as a compensatory anti-inflammatory mechanism dedicated to suppress unwanted immune-mediated tissue damage in the infected livers, and are also detectable in subjects with established chronic infection and in treatment non-responders^[23,25]. It has also been previously shown that there is an inverse correlation between IL10 production and the presence of HCV-specific T-cell proliferative responses in patients with established chronic HCV infection^[22,23], supporting the hypothesis that effective endogenous anti-inflammatory and immune inhibiting responses take place in association with a poorer outcome of the infection^[55].

The role of NK cells

Immunological events developing early on during the first phases of the acute infection are key in defining and shaping the subsequent adaptive responses. The innate immunological signatures in subjects with acute HCV infection differ in respect to their clinical outcome. Cytotoxic NK cells are proportionally predominant in comparison to CD56bright NK cells in acutely infected subjects who are spontaneously resolving the infection whilst this predominance is absent in patients progressing to chronicity^[24,56,57]. Furthermore, the relocation of innate and adaptive immune cells into the infected liver may be compromised in subjects progressing to chronicity, because these patients produce higher concentrations of CXCL10 antagonist in comparison to those who spontaneously resolve the infection as a consequence of a higher

dipeptidyl peptidase-4 (DPP4) activity present in their bloodstream^[24,58-61]. The role of DPP4 in skewing the clinical outcome of HCV infections has also been demonstrated anecdotally on the basis of these publications, in a case report of a patient with diabetes on the background of chronic HCV infection who was treated with the DPP4 inhibitor sitagliptin (a commonly used anti-diabetic drug) before undergoing antiviral treatment. During treatment with sitagliptin alone, the patient experienced a 2-log reduction in HCV viral load, which could not be ascribed to any cause other than the DPP4 inhibitor^[62].

DENDRITIC CELLS AND CHRONIC HCV INFECTION

DCs are professional antigen-presenting cells and have a primary role in the activation and coordination of primary immune responses. Different types of DCs exist through the body. They express different markers and home to different tissues; some re-circulate in the bloodstream^[3,63-65]. Dendritic cells sense pathogen-associated molecular patterns of bacterial, fungal or viral origin (PAMPs) thanks to their expression of several innate receptors such as TLRs and RLRs^[3,64]. Immature DCs are primarily sampling the immunological milieu of the tissue where they reside. However, upon activation immature DCs undergo a transformation process that includes upregulation of class I and class II MHC molecules and co-stimulatory molecules such as CD80 and CD86, production of type-1 and type-3 interferons (IFN α , IFN β , IFN λ), production of pro-inflammatory cytokines IL12, IL15, IL18 (amongst others), production of tissue-protecting IL10, and radical changes in their chemokine receptor and adhesion molecule profile^[3,63-65]. Activated mature DCs migrate to the lymphoid organs, where they interact with and activate both naive and experienced T cells. Interactions between DCs and NK cells are also important for the modulation of the innate immunity (reviewed in^[66]). The induction of a pro-inflammatory milieu locally also activates tissue-resident fibroblasts^[67-69], and the production of new extracellular matrix (fibrogenesis) marks the beginning of the healing response and the deposition of scar tissue. In the liver, this is associated with the activation of hepatic stellate cells (HSCs), which favours the development of fibrosis and cirrhosis, hallmarks of chronic liver disease^[70,71].

The idea that DC functions may be impaired during HCV infection has been evaluated in a multiplicity of studies, often with conflicting results depending on the experimental conditions and the cohorts of patients studied. However, on the overall, all these studies point at the non-equivocal conclusion that DCs do behave differently in patients with hepatitis C than they do in healthy subjects.

Liver dendritic cells in chronic HCV infection

As the liver is the primary site of HCV replication, it is conceivable that DCs would migrate into the infected tissues and that DC changes induced by the virus are prominent in the liver. Investigation of DCs in the liver of HCV-infected patients is indeed hampered by difficulties in gaining access to liver DCs. Therefore, our knowledge about liver DCs during HCV infection are limited. The few studies that investigated these cells demonstrated by either direct or indirect methods that DCs are enriched in the liver of HCV-infected patients^[72-75] (reviewed in^[76,77]). Contrasting results have been reported regarding the reciprocal distribution of DC subsets in the liver^[73,78]. Notably, the demonstration that DCs in livers from patients with hepatitis C localize in areas of liver necrosis^[79], together with the experimental observation that DCs and T cells are together recruited within HCV-infected human liver slices^[80], strongly support a role for DCs in the activation of HCV-specific immune responses. The mechanisms underlying DC enrichment in the liver of HCV-infected individuals are still poorly defined. Beyond an increased local recruitment, the observation that *in vitro* the migration of DCs is strongly inhibited by the interaction of DCs with the viral protein HCV E2^[81] may suggest that DC entrapment within the liver may contribute to the process.

Dendritic cells in the peripheral blood

pbDCs are the most accessible source of DCs. They can be divided into two main subsets: myeloid DCs (mDCs) and plasmacytoid DCs (pDCs)^[3,63-65]. Neither of them express lineage-specific markers (CD3, CD14, CD16, CD19, CD20), but both of them express high levels of HLA-DR. mDCs are characterized by high expression of the integrin CD11c and the blood DC antigens (BDCA) 1 (CD1c) or 3 (CD141). pDCs do not express CD11c and BDCA1-3, but express high levels of the IL3 receptor (CD123), BDCA2 (CD303) and BDCA4 (CD304) instead^[82]. Activated mDCs and pDCs have very distinct cytokine profiles. mDCs produce preferentially IL12 and IL10, whilst pDCs are the strongest producers of type-1 and type-3 interferons (IFN α , IFN λ)^[3,63-65]. pDCs express high levels of TLR3, 7-9, and are therefore highly sensitive to viral nucleic acids, nucleobases and ribonucleosides. mDCs may play a stronger role as orchestrators of pro-inflammatory responses, but pDCs are certainly strongly involved in the development of anti-viral responses. mDCs and pDCs can be counted and characterized by flow cytometry directly performed on peripheral blood samples^[83,84].

The frequency of pbDCs in the bloodstream is extremely low. This has hampered the study of these cell populations, as high concentrations of pbDCs can only be obtained starting from high volumes of peripheral blood. However, DCs can be successfully differentiated from peripheral blood monocytes

stimulated with either IL4 or IFN α in the presence of GM-CSF (monocyte-derived DCs, moDCs)^[85], and this has allowed researchers to bypass the issue of low availability in the peripheral blood and has granted them a useful *in vitro* model for the functional characterization of these cells.

Impairment of peripheral blood DCs in chronic HCV infection

Several studies investigated the impact of HCV infection on DCs by analyzing pbDCs. The main findings are illustrated in Figure 1, that recapitulates the current view on pbDCs involvement in HCV pathogenesis. In particular, most studies reported a numerical reduction of pbDCs in patients with chronic HCV infection, yet with some conflicting results on whether either mDCs or pDCs or both subsets are affected^[22,73,86-92]. Although pDCs are undoubtedly specialized in antiviral defenses, the reduction of mDCs may be relevant to HCV infection as well, as mDCs through IL12 production and the subsequent polarization of Th1 responses may contribute to the activation of cellular immunity. However, the reason for pbDC reduction is still unclear. pbDC reduction has been reported to be associated with the degree of liver inflammation^[93], possibly suggesting that the reduction of DCs in the peripheral blood may be due, at least in part, to an enhanced recruitment of these cells in the inflamed liver. pbDC reduction has also been reported to be more pronounced in patients infected with HCV genotype 2^[22], but not correlated with the viral load^[22,73,86,88,91], suggesting that multiple viral and non viral mechanisms may directly and indirectly contribute to the decrease of mDCs and pDCs in the circulation. Notably, our previous demonstration that the number of pbDCs is unaffected in healthy HCV-seropositive patients who underwent spontaneous resolution of their HCV infection^[22] clearly indicate that active HCV infection is needed in order to determine a reduction of pbDCs.

The functional assessment of pbDCs could also demonstrate that pbDCs from patients with chronic HCV infection are functionally impaired. Several studies demonstrated indeed that, upon stimulation with TLR ligands, viruses or interaction with T cells, both mDCs and pDCs from HCV-infected patients have impaired production of pro-inflammatory and anti-inflammatory cytokines compared with healthy controls. Due to the important role played by pDCs and IFN in antiviral immune responses, many Authors investigated the ability of pDCs to produce IFN α in response to distinct TLR ligands, with impaired IFN production observed in HCV-infected patients by some Authors^[87,88,90,91], but not by others^[89,94,95]. Moreover, mDCs from patients with chronic HCV infection were demonstrated to be characterized by impaired production of IL12, that in some cases was associated with impaired allostimulatory activity, and increased production

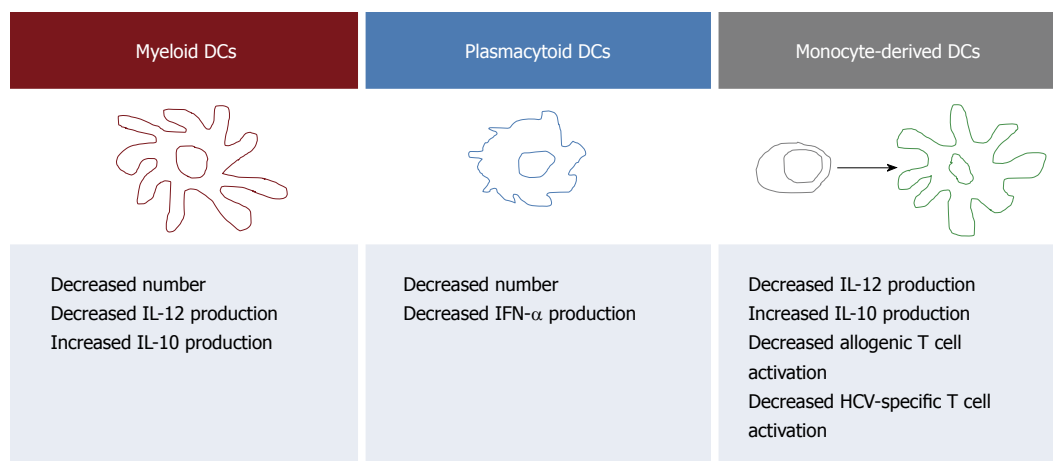


Figure 1 Summary of the hepatitis C virus-associated alterations of dendritic cells detectable in the peripheral blood. Patients with chronic hepatitis C undergo changes in the number and function of both myeloid dendritic cells (mDCs) and plasmacytoid DCs (pDCs) that may be relevant to the pathogenesis of chronic HCV infection. In particular, several Authors demonstrated that pDCs, which are specialized in antiviral immune responses, are decreased in HCV-infected patients and are endowed with a reduced ability to produce interferon (IFN). mDCs, that contribute to the activation of cellular immunity through the production of interleukin (IL)12, are decreased as well and characterized by reduced production of IL12 and increased production of the immunosuppressive cytokine IL10. These last functional defects are shared by monocyte-derived DCs that are also endowed with decreased ability to activate T cells. All references are reported in the text.

of IL10^[22,87,88,96,97]. All together, these observations indicate that, despite discrepancies among studies possibly due to differences in the patient cohorts and the experimental procedures, the analysis of circulating pbDCs can detect numerical and functional alterations of DCs occurring during HCV infection.

Similarly to pbDCs, also moDCs have been demonstrated to be impaired in HCV-infected patients, with reduced IL12 production, reduced allostimulatory activity and increased IL10 production reported in many studies but not confirmed in others, as reviewed elsewhere^[92]. moDCs have been suggested to be less representative of *in vivo* functions compared with pbDCs, as discussed elsewhere^[22]. Nevertheless, they have proven to be a very useful model for the study of DC functions. Yet, they have been largely used for studying the effects of cell culture-adapted strains of HCV on DCs and they provided insight into the molecular interactions of recombinant HCV proteins with DCs *in vitro* (reviewed in^[98]). Although the mechanisms whereby HCV affects DC function still remain largely elusive, the effects of many viral components and proteins on DCs have been defined^[99].

Effects of therapy with pegIFN plus RBV on peripheral blood DCs

As reported above, the analysis of pbDCs from patients with chronic HCV infection allows the detection of DC defects that can be relevant to the inadequate activation of immune responses leading to HCV persistence. Until recently the only therapeutic option for chronic HCV infection was the dual therapy with pegIFN plus RBV that achieved a sustained SVR only in part of the patients. For these reasons, several studies investigated the effects of dual therapy on pbDC number and functions according to virological

response, in order to define the impact of treatment on DCs and to possibly identify correlates of clinical response. In most of the studies, the standard treatment with pegIFN plus RBV for 24 or 48 wk induced an increase of the number of pDCs in patients who achieved a SVR^[100-102]. An early increase of pDCs during treatment was considered a correlate of favourable response in two studies^[100,103]. pDC increase was accompanied by mDC increase in one study^[102]. Only one study reported no variation in the number of pbDC subsets during treatment follow-up^[104], but the reasons for this discrepancy are not evident. In this context, we can add our evidence supporting the impact of the dual therapy on pDCs. We analyzed pbDCs in a cohort of 14 HCV-infected patients (11 males, 3 females; mean age 49 years, range 29-68 years) undergone standard treatment with pegIFN and RBV for 24 or 48 wk depending on the viral genotype. pbDCs were analyzed before, after 1 mo of treatment, at the end of therapy and 6 mo after treatment completion. pbDCs analysis was performed by flow cytometry as previously described^[22], in order to assess DC count, subset distribution, expression of activation/maturation markers (CD80, CD86, CD40, CD83), expression of molecules involved in HCV entry (CD81, DC-SIGN) and production of cytokines (IL12 and IL10). As shown in Figure 2, we observed that pegIFN and RBV treatment restored correct proportions of pDCs in SVR patients by inducing a rapid and progressive increase of these cells that was observed after 1 mo of treatment, further increased at the end of treatment and persisted after the end of treatment. SVR patients showed higher pDC levels than non-SVR at any time point. pegIFN and RBV treatment also induced a transient increase of IL12 production by pbDCs, that was observed after 1 mo of treatment but decreased thereafter, and that was

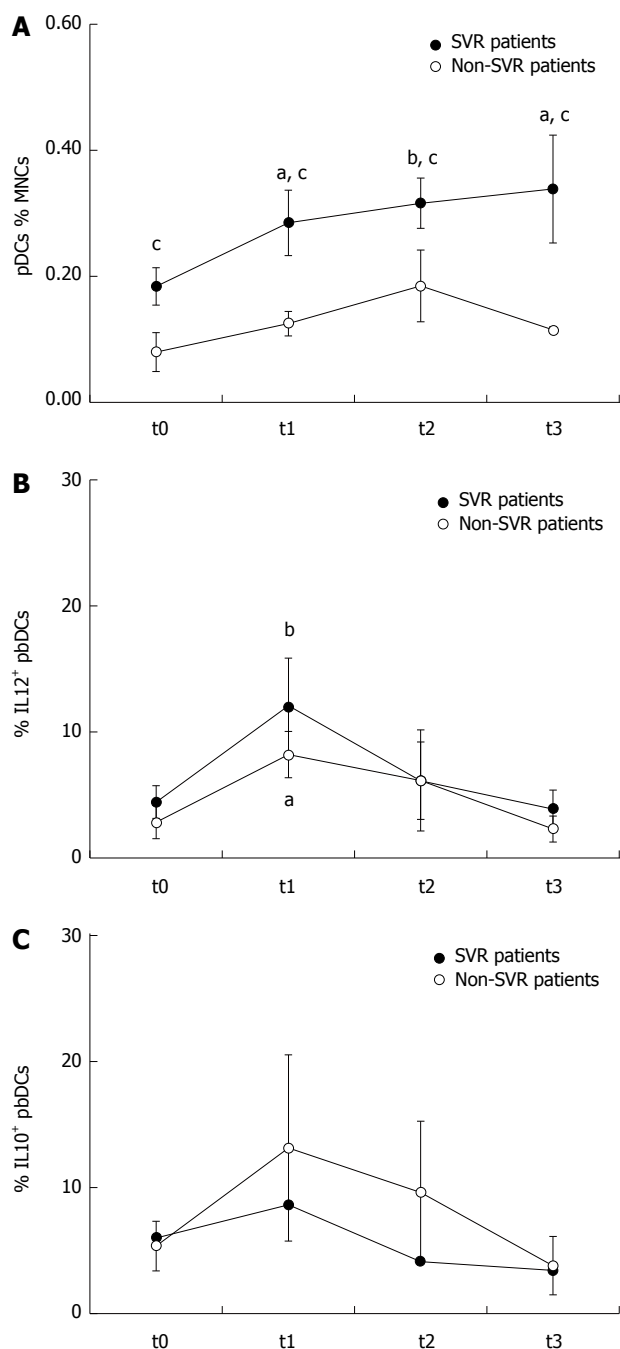


Figure 2 Effects of dual therapy with pegylated interferon and ribavirin on peripheral blood dendritic cells. The treatment induced a rapid, progressive and persistent increase of plasmacytoid dendritic cells (pDCs) in sustained virological response (SVR) but not in non-SVR patients. The frequency of pDCs was higher in SVR than non-SVR patients at all the time points. pegIFN and RBV treatment also induced a transient increase of peripheral blood DC production of interleukin (IL)12, that was observed after 1 mo of treatment but decreased thereafter, and that was more pronounced in SVR than non-SVR patients. IL10 production showed a similar trend and tended to be higher in non-SVR patients. a and b: $P < 0.05$ and $P < 0.01$, respectively, any time vs t0 within each group, as assessed by Wilcoxon signed rank test; c: $P < 0.05$, SVR vs non-SVR, as assessed by Mann-Whitney test. t0: Before treatment; t1: After 1 mo of treatment; t2: At the end of treatment; t3: 6 mo after treatment completion.

more pronounced in SVR than non-SVR patients. The other pbDC parameters were not affected by the dual therapy. Because of the role played by chemokines

and chemokine receptors in regulating DC migration, Mengshol and coll. also investigated the expression of CXCR3 and CXCR4 demonstrating that these molecules, upregulated on pbDCs at baseline, are normalized by pegIFN plus RBV, as well^[101].

The effects of therapy have also been assessed on moDCs, in order to investigate the mechanisms responsible for the impaired stimulation of T cells by DCs from HCV-infected patients. In an elegant cross-over study involving chronic HCV patients undergoing anti-viral treatment with pegIFN and RBV^[105], peripheral blood DCs and T cells were independently purified from the same patient at two time points: baseline (BL), with high serum HCV viral load, and treatment week 12 (TW12), with pharmacologically reduced serum HCV viral load. Co-cultures of autologous T cells and DCs were prepared using all four possible combinations from each subject (BL DCs plus BL T cells or TW12 T cells, and TW12 DCs plus BL T cells or TW12 T cells) and DC-induced T cell responses were measured in order to ascertain the role of DCs or T cells in the context of altered HCV-specific responses. The study showed that the measurable reduction in T-cell responses was not dependent on the T cells used, but it was always only the use of DCs obtained at baseline, with high viral presence, which was associated with low T cell immunity. Conversely, DCs obtained at the later time point, after pharmacological viral clearance, were able to stimulate with similar higher efficiency both BL T cells and TW12 T cells.

Peripheral blood DCs as predictors of response to treatment

Predicting the outcome of a treatment that is administered to a patient is a goal for both patients and physicians. Accordingly, the identification of pbDC parameters measured at baseline or at the beginning of treatment that may predict SVR or non-SVR was a primary aim of the studies that analyzed pbDCs in HCV-infected patients treated with pegIFN and RBV. Notably pDCs, whose increase represented the most relevant treatment-induced pbDC modification, also resulted good predictors of response in two studies, where higher levels of pDCs during the first weeks of treatment were associated with SVR^[100,103]. High levels of pDCs at baseline was a predictor of SVR in our cohort of patients, as well, independent from other clinical features including viral genotype and viral load (Figure 2). An increased pbDC production of IL-10 at baseline has been reported as non-SVR predictor by Liang and coll^[102], and confirmed in our cohort of patients.

CONCLUSION

Although the impact of HCV on DC phenotype and function is likely to be more prominent in the liver as it is primarily infected by the virus, nevertheless several HCV-induced alterations are detectable in

pbDCs that represent the most accessible source of DCs. These alterations include numerical reduction, impaired production of IL12 and IFN α , impaired allostimulatory activity and increased production of immunosuppressive IL10. Despite somehow controversial because variably observed among different cohorts of patients, these HCV-associated DC features are relevant to our understanding the immune mechanisms underlying the propensity of HCV to establish persistent infection. Importantly, the non-invasive accessibility of pbDCs renders the analysis of these cells a convenient procedure that can be serially repeated in patient follow-up. Accordingly, the study of pbDCs in HCV-infected patients during conventional treatment with pegIFN and RBV indicated that restoration of normal pDC count may represent a further novel mechanism contributing to the efficacy of the dual therapy. It also identified pDCs and IL10 as putative predictors of response to therapy. Treatment of chronic HCV infection is changing, as new generation direct-acting antivirals will soon be available for use in interferon-free therapeutic strategies. The phenotypic and functional analysis of pbDCs in this novel therapeutic setting will provide a valuable tool for investigating mechanisms underlying treatment efficacy and for identifying predictors of treatment response.

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