

Immunological scars after cure of hepatitis C virus infection: Long-HepC?

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Hepatitis C virus (HCV) infection provides a unique opportunity to study the effects of spontaneous or treatment-induced viral elimination on the human immune system. Twenty to 50% of patients with acute HCV infection spontaneously clear the virus, which is related to the quality of the individual's immune response, while the chronic infection is associated with an altered and impaired immune response. Direct-acting antiviral agents are now available that provide sustained viral elimination in more than 95% of patients with chronic HCV infection. Viral elimination leads to a decrease in disease sequelae such as cirrhosis and hepatocellular carcinoma, and extrahepatic manifestations also improve. However, some patients may still experience long-term complications, and viral elimination does not protect against HCV reinfection. This review addresses the question of whether the altered and impaired immune response caused by HCV normalizes after viral elimination and if this may affect the long-term clinical course after HCV cure.

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Introduction

Hepatitis C virus (HCV) infections remain a major global health burden, with 1.5 million new HCV infections per year and 58 million chronic HCV infections worldwide. HCV infection either leads to acute, resolving hepatitis or develops into chronic hepatitis with the long-term risk of developing cirrhosis and hepatocellular carcinoma (HCC) [1]. In addition, HCV infection is associated with various extrahepatic manifestations such as fatigue, insulin resistance, or cryoglobulinemia. Chronic HCV infection is associated with an altered and impaired immune response (reviewed in Ref. [2]), which may explain the high rate of chronic infections, but also the risk of reinfection and, in part, the long-term consequences of HCV infection.

Fortunately, interferon (IFN)-free therapies with direct-acting antiviral agents (DAA) have been available for several years. DAAs target either the HCV protease, the HCV protein NS5A, or the HCV polymerase. A combination of an NS5A inhibitor with a protease or polymerase inhibitor is first-line therapy, and if treatment fails, re-treatment with a combination of all three DAA classes is possible. Overall, DAA treatment is well tolerated and now results in a sustained virologic response rate (SVR) of well over 95%, leading to a substantial reduction in liver morbidity and mortality [3]. However, SVR does not protect against reinfection which is of extreme importance, especially for individuals with high risk behaviors [4], and not all sequelae of chronic hepatitis C (CHC) appear to be completely reversible after SVR. Patients with advanced fibrosis or cirrhosis have a residual risk of HCC [5] and impaired quality of life or extrahepatic manifestations are only partially improved or are not reversible in all patients (reviewed in Ref. [3]).

An important question, therefore, arises as to whether the altered immune response normalizes after viral elimination and whether this has an impact on the long-term clinical outcome of patients.

Finally, in contrast to HIV or HBV infections, where complete viral elimination is not or very rarely possible, HCV infection provides a unique model to study the effects of complete viral elimination on changes in immune responses. This may lead to a conceptual understanding of how impaired immune responses can be

restored, which is important not only for chronic infections but also for cancer. In the following sections, we review our (Table 1) and other studies on the immunologic effects of HCV elimination by DAA therapy and the possible consequences for the long-term outcome (Figure 1).

Effects of hepatitis C virus and its elimination on the systemic inflammatory environment

HCV infection results in rapid induction of IFN-stimulated genes (ISG), which elicits a strong type I/III IFN response. Despite this IFN response, most patients with acute HCV infection are unable to completely clear the virus and progress to chronic infection due to various evasion mechanisms. Because of viral persistence, ISG induction and the type I/III IFN response persist and correlate with viral load (reviewed by Ref. [2]). Besides type I/III cytokines also other cytokines and chemokines are altered in patients with acute and chronic HCV infection [6–8]. This inflammatory milieu in HCV infection may also affect the overall immune responses. For example, we have shown that CMV- and EBV-specific T-cell responses in patients with chronic HCV infection have distinct profiles with higher expression of co-regulatory receptors (e.g. PD-1 and Tim-3) that could be associated with type I IFN by demonstrating that IFN alpha stimulation induced further upregulation of these markers *in-vitro* [7]. The type I/III milieu may also help to explain clinical features of HCV infection such as chronic fatigue. We have demonstrated that plasma levels of soluble inflammatory mediators (SIM), especially IFN alpha and CXCL-10 but also distinct activation phenotypes of immune cells were associated with neuropsychological parameters [9].

Treatment with DAA and subsequent HCV elimination can potentially normalize the HCV-related inflammatory milieu, for example by downregulation of ISG [10]. However, in the short-term follow-up (FU) of up to 24 weeks after the end of therapy, there does not appear to be complete recovery. We have reported that upregulated SIM levels such as CXCL-10 decreased during and after successful therapy, but not to normal levels in all patients, and that suppressed parameters such as IL-17 did not increase [8]. Similar observations were made in individuals who cleared HCV after liver transplantation [11].

The long duration of HCV exposure of sometimes more than decades in CHC may be one reason for the long-lasting imprinting of the inflammatory milieu after viral elimination. Indeed, a study by Rosenberg et al. showed that gene expression profiles in peripheral blood rapidly normalized to pre-infection levels after spontaneous resolution of acute HCV infection in patients who had thus been viremic for less than six months. However, ISG expression interestingly remained altered over a longer

period of time compared with samples from acute dengue virus infection [12]. We have also studied the soluble inflammatory milieu in patients with acute HCV infection [6] who cleared HCV upon DAA therapy in a prospective study [13]. Similar to the study in chronic patients, the majority of SIMs decreased, but not all normalized, and SIMs that were suppressed before DAA treatment remained downregulated [6]. Thus, even after acute HCV infection, the inflammatory response does not seem to completely normalize. However, so far most of the studied patients after both chronic and acute HCV infection were followed up only 12–24 weeks after the end of treatment (EOT). Studies with longer FU times are needed to also analyze the effect after prolonged virus elimination. In addition, the impact of the cytokine milieu after HCV elimination on liver disease progression and extrahepatic manifestations such as chronic fatigue remains to be further determined. Published data suggest a link between systemic inflammation and progression of liver disease and liver cancer in patients with advanced chronic liver disease [14]. We have demonstrated that distinct SIM were associated with the development of HCC after HCV elimination by DAA therapy and this SIM plasma profile of cirrhotic patients who developed HCC after therapy, in contrast to those who remained HCC-free [15].

Likewise, the effects of HCV elimination on the immune response to other pathogens are of interest. The changes in the ISG and inflammatory milieu may influence the heterologous immune response. For example, we have shown that the dominance of HCV and higher serum interferon- γ -induced protein 10 (or CXCL-10) levels was associated with lower quantitative hepatitis B surface antigen in HBV/HCV-coinfected patients [16]. Interestingly, reactivations of hepatitis B virus [17] but also herpes virus [18] infections have been documented after DAA therapy, suggesting that indeed modulations of immune responses after HCV elimination may have clinical implications. The hypothesis as to why reactivation of viral infections occurs may be due to the fact that interferon responses are key elements of antiviral defense, and these are down-regulated after HCV eradication [19]. On the other hand, the reduced inflammatory environment may also affect cellular immune responses to other pathogens. We have shown that PD-1 expression on CMV- and EBV-specific CD8⁺ T-cells decreases after DAA therapy and HCV elimination [7]. Furthermore, direct antigen-specific effects, for example, through T-cell cross-reactivity [20] might also contribute to changes in T-cell responses to other pathogens.

Effects of hepatitis C virus and its elimination on epigenetic imprints

Due to persist antigen stimulation T-cells in chronic infections develop into an exhausted T-cell state.

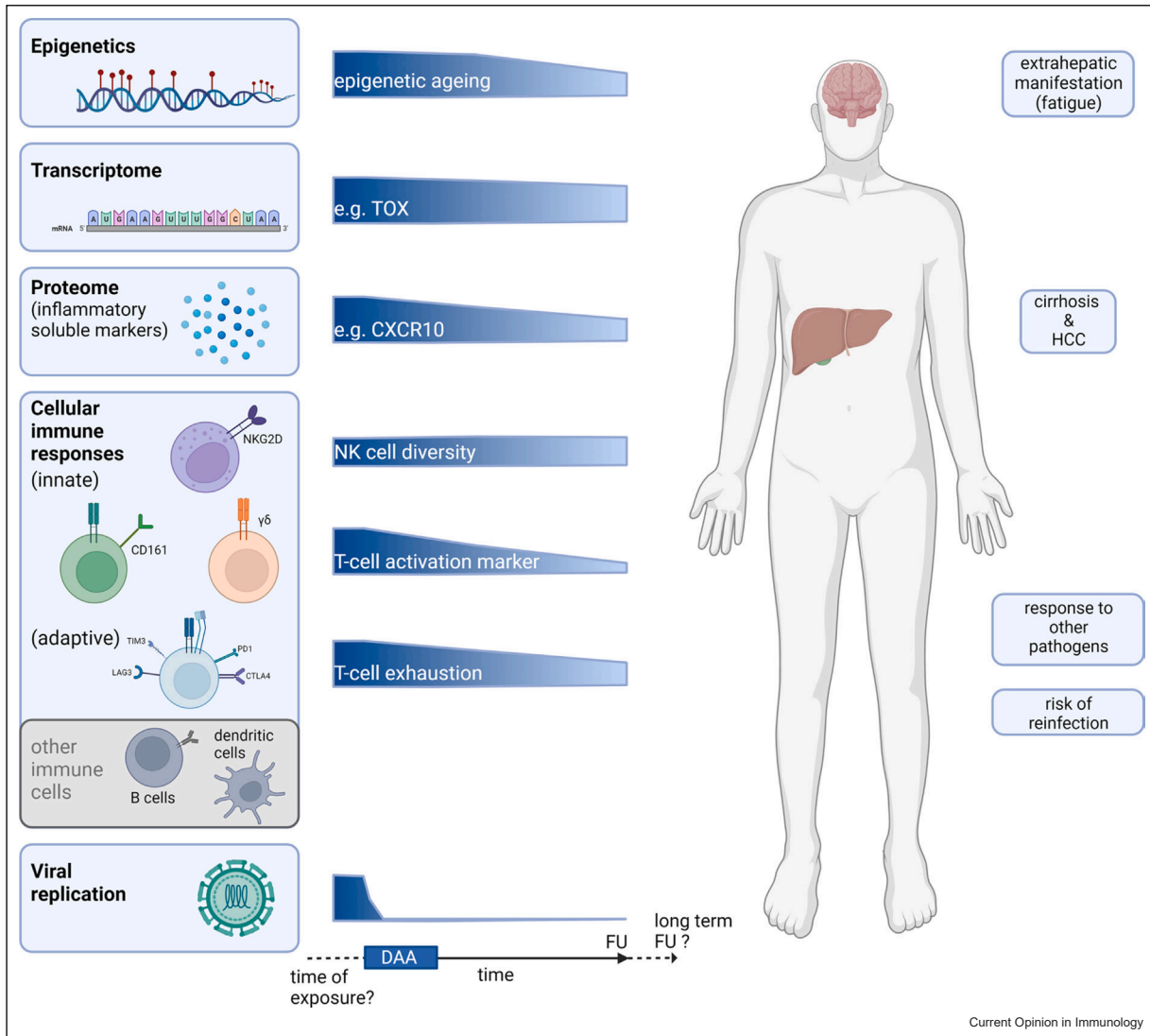
Table 1

Studies conducted in the CRC900 consortium on acute and chronic HCV infection before and after HCV elimination by DAA therapy.

Ref.	Cohorts	Main results
Inflammatory milieu (Proteomics (SIM))		
[7]	CHC patients (n = 76, 69 without, 7 with DAA), healthy controls (n = 78)	In CHC patients, plasma concentrations of cytokines, especially type I/III IFN, were altered. DAA therapy significantly decreased IFN α levels.
[8]	CHC patients (fibrosis and cirrhosis) treated with DAA, FU 12–24 weeks after EOT (n = 28), healthy controls (n = 5), patients with steatosis (n = 20)	The expression patterns of cytokines and chemokines in patients with chronic HCV infection were altered compared with controls. The majority of upregulated analytes decreased during and after DAA therapy, but HCV elimination did not result in complete restoration.
[9]	Patients with chronic liver disease with or without neuropsychiatric symptoms (HCV n = 14, other chronic liver diseases n = 26)	In patients with chronic inflammatory liver disease and neuropsychiatric symptoms, SIM were significantly more upregulated, especially IFN α and CXCL-10, than in patients without symptoms. HCV patients showed a unique pattern of immune alterations.
[11]	Liver transplant recipients (LT): HCV+LT+DAA (n = 25, FU 12–24 weeks after EOT), non-HCV+LT (n = 14), healthy controls (n = 10)	In patients cured of HCV by DAA after liver transplantation, the altered inflammatory milieu did not normalize, nor did the subsets of peripheral immune cells normalize to the levels seen in the absence of HCV recurrence.
[15]	CHC patients with cirrhosis (n = 31, 15 developed HCC, 16 without HCC), healthy controls (n = 10)	Elevated levels of SIMs pre DAA therapy in cirrhosis patients were associated with HCC development post DAA therapy.
[6]	Acute HCV (n = 20, with DAA therapy, FU 24 weeks), chronic HCV (n = 23, with DAA therapy), healthy controls (n = 20)	Profound SIM changes were observed in acute HCV patients. During DAA treatment and FU, the majority of SIM decreased, but not all normalized.
Epigenetics		
[28]	CHC without cirrhosis (n = 22), CHC with cirrhosis (n = 24), CHC with cirrhosis + HCC (n = 8), all treated with DAA therapy, FU up to 96 weeks after EOT	Chronically HCV infected patients showed a significant increase in overall DNA methylation and an increase in epigenetic age acceleration. Elimination of HCV partially reversed the accelerated epigenetic ageing in long-term FU; however, patients with HCC show no change in the acceleration of epigenetic ageing.
Innate cellular immune responses		
[35]	CHC patients (n = 26–42, treated with DAA: FU 12–24 weeks after EOT (n = 26), FU up to 96 weeks after EOT (n = 21)), healthy controls (n = 10–22)	Chronic HCV infection induced a functional imprinting on human NK cells, especially diversity. This imprinting was largely irreversible and remained long after the virus was eliminated.
[37]	CHC patients (n = 23, treated with DAA), healthy controls (n = 9)	Clonality and complexity of the $\gamma\delta$ -TCR repertoire were largely comparable to those of healthy controls with more V γ 9 ⁺ T-cells and an increase in V δ 3 clones. Only minor changes in $\gamma\delta$ -TCR repertoires were observed after DAA therapy.
[38]	CHC patients (n = 28, treated with DAA), controls with fatty liver disease (n = 20)	In CHC, the frequency, phenotype, and function of MAIT cells in blood were altered. After DAA therapy, MAIT cells were not restored.
[44]	Acute HCV (n = 15 with DAA therapy, FU 24 weeks after EOT), chronic HCV (n = 12, with DAA therapy, FU 48 weeks after EOT), healthy controls (n = 14)	The markedly activated phenotype of unconventional T-cells (e.g. $\gamma\delta$ -T cells) during acute HCV infection was reversible after early DAA treatment. However, MAIT cell dysfunction was not restored.
Adaptive cellular immune responses		
[47]	CHC patients without DAA therapy (n = 86), acute HBV (n = 6), acute HCV (n = 13), healthy controls (n = 49)	The study assessed 2B4 expression on HCV-specific CD8 ⁺ T-cells during acute and CHC. 2B4 cross-linking could lead to both inhibition and activation of HCV-specific CD8 ⁺ T-cell function, depending on expression levels of 2B4 and the intracellular adaptor molecule SAP.
[46]	CHC patients (n = 51, without DAA therapy)	HCV-specific exhausted CD8 ⁺ T-cells were characterized by an individualized hierarchy of coregulatory receptor (PD-1, CTLA-4, TIM-3, or 2B4) usage.
[7]	CHC patients (n = 76, 69 without, 7 with DAA), healthy controls (n = 78)	DAA-induced HCV elimination partially reversed the phenotype (higher PD-1 expression) of CMV/EBV-specific CD8 ⁺ T-cells in patients with CHC.
[49]	CHC patients (n = 47, n = 28 without cirrhosis, n = 12 with cirrhosis; all treated with DAA, FU 24 weeks after EOT), healthy controls (n = 7)	HCV-specific CD8 ⁺ T-cell phenotypes and functional responses were only partially restored after DAA-induced HCV clearance in CHC. The impaired mitochondrial fitness of HCV-specific CD8 ⁺ T-cells was not altered after HCV elimination.

DAA – direct-acting antiviral agents; FU – follow-up, EOT – end of treatment, SIM – soluble inflammatory mediators; HCC – Hepatocellular carcinoma.

Figure 1



Schematic illustration of immunological alterations as a result of HCV elimination by DAA therapy in patients with chronic HCV infection (created with BioRender.com).

Recent studies have also shown an impact on the epigenetics of these cells imprinting the exhausted phenotype [21,22]. Chromatin accessibility analysis of HCV specific CD8⁺ T-cell responses reveals epigenetic signatures of exhaustion that appear to be reversible only to a limited extent after HCV elimination, for example, an epigenetic scar remains in super-enhancers near the *TOX* (thymocyte selection-associated HMG BOX) and *HIF1α* (hypoxia inducible factor-1α) genes [23]. *TOX* is an important regulator of T cell exhaustion [24] and *HIF1α* is a transcriptional regulator of responses to hypoxia and has been linked to T cell activation [25]. Thus, epigenetic scars at loci near critical transcription factors associated with T cell responses may help explain the maintained exhausted immune responses [23].

In addition to showing that the detected epigenetic imprints are not fully reversible after SVR, two groups (Hamdane et al. and Hlady et al.) have detected genome-wide changes in the acetylation of histones (such as H3K4me3, H3K4me1, H3K27ac, and H3K27me3) in hepatocytes that have been linked to increased liver cancer risk [26,27].

In our recent study, we analyzed the epigenetics of immune cells from chronically HCV-infected patients based on DNA methylation. We found a different epigenetic profile in chronically HCV-infected patients compared with healthy controls. These differences could also be reflected by calculating epigenetic age acceleration according to the Horvath clock, patients with CHC

are biologically older. After cessation of DAA therapy (8–12 weeks), epigenetic age acceleration did not improve. Nevertheless, at long-term FU 96 weeks after the end of therapy, we were able to show a significant reduction in age acceleration, which, however, was not completely normalized to the level of healthy controls. Interestingly, patients who developed HCC after DAA therapy showed no improvement in their biological age [28].

The question remains whether the infection with HCV itself or the above-mentioned inflammation is the main trigger for the epigenetic modifications. This needs further investigation, also to discover a treatment that reverses the epigenetic imprints.

Effects of hepatitis C virus and its elimination on cytotoxic cellular innate immune responses

Natural killer (NK) cells are highly enriched in the human liver and contribute as innate lymphocytes to antiviral immune responses. Many studies have suggested an important role for NK cells in the pathogenesis and control of HCV infections (reviewed in Ref. [29]). In chronic HCV infection, NK cells have impaired functions and an altered phenotype [30], which is attributed to the persistent exposure to HCV-induced type I IFN responses [31]. DAA-mediated HCV elimination has been associated decrease in type I IFN responses and, accordingly, normalization of NK cell phenotype and functions [32–34]. However, we have shown that some imprints on NK cells caused by chronic HCV infection do not appear to be completely reversible despite viral elimination. In detail, long term chronic HCV infection increased the inter-individual, but decreased intra-individual NK cell diversity, which was not reversed after DAA therapy and HCV clearance [35]. We are currently investigating the NK cell response in patients with acute HCV infection to determine if a shorter exposure to HCV makes a difference.

Unconventional T-cells, such as $\gamma\delta$ T-cells and Mucosal-associated invariant T (MAIT) cells are unique innate-like T-cells that bridge innate and adaptive immunity. It has been shown that $\gamma\delta$ T-cells are less efficient in producing cytokines and exhibit an activated phenotype in chronic HCV infection [36]. After DAA therapy and clearance of HCV, the $\gamma\delta$ T-cells lose their activated phenotype, but stay dysfunctional [36]. We have investigated if the alteration of the $\gamma\delta$ T-cell compartment is due to skewing in the T-cell receptor repertoire, but this remains stable even after HCV cure with DAA therapy [37].

It was also shown that dysregulation of MAIT cells might play a role in the progression of chronic HCV

infection. We and others have shown that these dysfunctions cannot be restored or can only be partially restored after HCV cure achieved by DAA therapy [38–41]. However, the degree of liver inflammation and liver fibrosis may contribute to the degree of dysfunction, loss, and recovery of MAIT cells [40,41]. Alterations in unconventional T-cells in HCV patients and after HCV cure may be important for patients with cirrhosis and especially cirrhosis-related immune dysfunction, as these cells may contribute to immunopathogenesis and immune protection in this setting [42,43].

We investigated whether brief HCV exposure in the setting of acute hepatitis C has an impact on the phenotype and function of unconventional T-cells. We investigated whether a short HCV exposure in the setting of acute hepatitis C has an impact on the phenotype and function of unconventional T-cells. However, while the phenotype of $\gamma\delta$ T-cells and MAIT cells normalized their activated phenotype, the frequency and function of MAIT cells remained impaired compared with healthy controls up to 24 weeks after viral elimination by DAA therapy [44].

Effects of hepatitis C virus and its elimination on cellular adaptive immune responses

CD8⁺ and CD4⁺ T-cells are the main contributors to the cellular adaptive immune response. A strong and broad virus-specific T-cell response is associated with spontaneous HCV elimination, whereas failure of the T-cell response, either by viral escape or T-cell exhaustion, is associated with the development of chronic infection (reviewed by Refs. [2,45]). Exhaustion of the HCV-specific T-cell response is a hallmark of chronic HCV infection and is characterized by increased coexpression of inhibitory receptors (e.g. PD-1, Tim-3, CTLA-4, and Lag-3, 2B4) [46,47], distinct epigenetic [23], transcriptional [48], and metabolic [49] alterations compared with effector or memory T-cells and impaired functional response (reviewed by Refs. [2,45]). T-cell exhaustion is driven by high and persistent antigen stimulation and in this context, it is a mechanism to prevent an overwhelming immunopathology [50]. The question is whether elimination of the antigen can restore exhausted T-cell responses, which can be addressed using HCV infection as a human model system.

Martin et al.'s study was the first to show an increase in HCV-specific CD8⁺ T-cells after DAA therapy with a recovery of CD8⁺ T-cell proliferative capacity [51]. However, further studies, including our own, showed only partial recovery of the phenotype of HCV-specific T-cells and even less restoration of the functional response [49,52,53]. For example, we have shown that mitochondrial dysfunction of exhausted HCV-specific CD8⁺ T-cells

was not restored in terms of mitochondrial polarization, mitochondrial mass, and reactive oxygen species back to the level of healthy donors [49]. These data indicate that during chronic HCV infection cells develop dysfunctions, which cannot be completely reversed by antigen removal. Recent studies support this finding of a molecular scar also on a transcriptomic level [48]. Furthermore, this concept is supported by the finding that expression of the central transcription factor *TOX*, which controls T-cell exhaustion, decreases after antigen removal but remains higher than in other memory T-cells after recovery from chronic HCV by DAA therapy [24,52]. Importantly, T-cells from chronic HCV infection that were exposed to antigen for a limited time because of the emergence of viral escape mutations were functionally and transcriptionally more similar to memory T-cells from spontaneously resolved HCV infection [52]. In line with this, in one study, HCV-specific T-cell responses were stronger after DAA treatment of acute HCV infection than after DAA treatment of CHC and were similar to those found after spontaneous HCV elimination, except that the breadth of the T-cell response was greater with spontaneous HCV elimination [54].

The before mentioned epigenetic scarring also translates into a molecular scar on the transcriptomic level [23]. This phenomenon of immunological scarring is not only restricted to chronic HCV infections but has also been found in other chronic infections. For example, in chronic lymphocytic choriomeningitis virus (LCMV) infection, similar immunological scars have been identified with functional impairment of virus-specific T cells [55,56]. In addition to this functional impairments, also epigenetic scars have been described to have an impact on memory formation and stay also after clearance of chronic LCMV [23,57].

This highlights, that immunological scarring might be from importance in many chronic viral infections and a possible new therapeutic target.

Most of the data described above were obtained for CD8⁺ T-cells, but also HCV-specific CD4⁺ T-cells remained dysfunctional and in very low abundance after HCV cure by DAA therapy for CHC. This study showed comparable phenotypes and functions of CD4⁺ T-cells resembling protective memory after HCV vaccination and spontaneous HCV elimination [58].

Thus, the current data suggest that the duration of antigen exposure may be relevant for the potential of reversion of the impaired immune response.

In addition, other adaptive immune cells such as regulatory T-cells [59], B-cells [60], or dendritic cells [61] have been studied in the context of chronic HCV infection and cure. However, details will not be further discussed in the review.

Relevance for translation

To further improve the therapy of chronic viral infections such as chronic hepatitis B or HIV (no longer relevant for HCV as being curable), it is important to explore the reasons for immunological scarring and find out how to overcome it. As suggested above, epigenetic changes due to the modulated cytokine environment or the viral infection itself could be the reason for the scarred transcriptome profile and the impaired and metabolically altered T-cell response. To overcome this, correction of the deregulated metabolic pathways underlying T-cell dysfunction, as discussed in the context of hepatitis B virus infection, may represent a novel strategy [62,63]. Another approach could be therapy with epigenetic modifiers such as DNA methyltransferase inhibitors (e.g. 5-aza-2-deoxycytidine or decitabine) or histone deacetylase inhibitors, which in combination with checkpoint inhibition were able to restore an exhausted T-cell response in the LCMV mouse model [64].

The data might also be relevant to understand and also combat long term consequences of HCV infection after cure such as HCC but also extrahepatic manifestations [5]. Chronic fatigue, which is frequently present in patients with CHC is not always reversed after HCV cure [65]. Finally, targeting the impaired immune responses post HCV cure could potentially decrease the risk HCV reinfection.

Another open question that still arises is whether the HCV elimination influences the response to unrelated virus-specific [7] or even tumor associated [66] T-cell responses or if the scarring which can be found in the HCV-specific T-cells after HCV clearance can also be found in immune responses against unrelated antigens.

Finally, the discussion about a scar in the immune response is important for considering controlled human infections models that are considered to study HCV vaccines [67].

Summary and conclusion

In summary, improvement in certain features of the immune system can be observed after successful HCV treatment, but many features of immune exhaustion persist over time after the elimination of the virus. These data could explain the long-term consequences of infections as now discussed in the case of Long-COVID, and understanding the causative mechanisms could contribute to new therapeutic approaches or diagnostics.

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Data Availability

No data were used for the research described in the article.

Declaration of Competing Interest

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