Viral (hepatitis C virus, hepatitis B virus, HIV) persistence and immune homeostasis

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Summary

Immune homeostasis is a host characteristic that maintains biological balance within a host. Humans have evolved many host defence mechanisms that ensure the survival of individuals upon encountering a pathogenic infection, with recovery or persistence from a viral infection being determined by both viral factors and host immunity. Chronic viral infections, such as hepatitis B virus, hepatitis C virus and HIV, often result in chronic fluctuating viraemia in the face of host cellular and humoral immune responses, which are dysregulated by multi-faceted mechanisms that are incompletely understood. This review attempts to illuminate the mechanisms involved in this process, focusing on immune homeostasis in the setting of persistent viral infection from the aspects of host defence mechanism, including interferon-stimulated genes, apolipoprotein B mRNA editing enzyme catalytic polypeptide 3 (APOBEC3), autophagy and interactions of various immune cells, cytokines and regulatory molecules.

Keywords: hepatitis C virus; HIV; immune homeostasis; viral persistence.

Introduction

Immune homeostasis is a host characteristic that maintains biological balance. Humans have evolved many host defence mechanisms that ensure the survival of individuals upon encountering a pathogenic infection. Although healthy individuals coexist in equilibrium with the microbial world, disease can be viewed as a progression away from this immune homeostasis. Microorganisms disturb immune homeostasis because they invade and replicate, causing inflammation and destruction of the tissue, as well as depletion of the host's nutrients, ultimately leading to a chronic infection or death of the host if the infection is not resolved. A well-described example is intestinal homeostasis, which depends on complex interactions between microbes, the intestinal epithelium and the host immune system. Diverse regulatory mechanisms cooperate to maintain intestinal homeostasis, and a breakdown in these pathways may precipitate the chronic inflammatory pathology found in inflammatory bowel disease.¹ Hence, a balanced interaction between microbes and the host immune system is required to maintain health.²

In relation to blood-borne viruses that often result in chronic infections, most notably hepatitis C virus (HCV), hepatitis B virus (HBV) and the HIV, their survival in the host may also be a function of immune homeostasis. At the onset of infection, overly rapid viral replication leads to cell lysis, tissue injury and premature host death whereas inadequate viral replication may result in viral latency or clearance (Fig. 1). The virus must balance its own growth with the death of host cells and circumvent the immune response to survive. Strategies for viral survival include regulating apoptosis, inhibiting interferon production, modulating the MHC class I/II functions that ultimately affect the cytotoxic lymphocyte and natural killer (NK) responses, and limiting cytokine or chemokine production/function. Long-term viral survival (i.e. viral persistence or latency) requires down-regulation of lytic gene expression, inhibition of apoptosis and minimization of the inflammatory response. In other words, viral persistence requires host immune homeostasis.

Viral factors for persistent infection

Viruses persist in cells because they are able to down-regulate key processes that if left unattended would result in cell death. Specifically, regulation of viral transcription and genomic replication allows for long-term viral persistence. Many viruses (such as HCV, HBV and HIV) that cause persistent infection and chronic disease are successful at in vivo survival because of their cell tropism and

Figure 1. Viral persistence and immune homeostasis. With high viral load infection, virus may evade host immunity, leading to death (a) or illness (b) of the host. With anti-viral treatment, viral load may be reduced, leading to viral clearance or latency. Meanwhile, anti-viral treatment may cause mutation in the virus and drug-resistance (b). When immune homeostasis is maintained, the host exhibits asymptomatic persistent infection, whereby the immune system fight against the virus; and viral load fluctuates during chronic infection (c). Under a strong host immune response, viral infection may become latent (d) or be cleared (e).

ability to auto-regulate their replication efficiently within target cells. Common features of auto-regulation include sensors to the external environment, negative feedback loops, transcriptional enhancers and transcriptional silencers. In some cases, auto-regulation results in steady-state levels of virus replication; in other infections, the virus enters latency only to reactivate intermittently. Viruses have therefore developed numerous strategies for subverting the host defences. The biological characteristics of HCV, HIV and HBV are summarized in Table 1, and the viral factors for persistent infection are described below.

Hepatitis C virus

Binding and entry of HCV is believed to be a multistep process involving HCV envelope glycoproteins E1 and E2 as well as several attachment and entry factors such as CD81, scavenger receptor class B type I, members of the claudin family and occludin. $3,4$ This process involves several viral and host factors and is targeted by host-neutralizing responses. As the HCV envelope glycoproteins E1 and E2 interact with host cell factors and trigger the conformational changes necessary to initiate infection, they are important targets for virus neutralization. A recent study⁵ indicated that altered use of the CD81 receptor allowed the virus to escape neutralizing antibodies, and this altered use of CD81 is mediated by residues at positions 447, 458 and 478. Kinetic studies showed that these mutations affect virus–antibody interactions during the post-binding steps of the HCV entry process. Functional

studies with a large panel of patient-derived antibodies showed that this mechanism mediates viral escape, leading to persistent infection in general. Another study revealed that mutations in HCV E2 located outside the CD81 binding sites lead to escape from broadly neutralizing antibodies but compromise virus infectivity.⁶ Recently, it was reported that HCV has evolved mechanisms that antagonize host cell death signals so that viral propagation can continue unabated in infected cells. For example, HCV core protein blocks tumour necrosis factor- α (TNF- α) -mediated apoptosis signalling and inhibits caspase-8 activation.⁷ A review of how HCV interferes with the different steps of initial antiviral host-response and establishes persistent infections was summarized by Bode *et al.*⁸ HCV targets the Toll-like receptor 3 (TLR3), retinoic acid-inducible gene I (RIG-I), and interferon (IFN) signalling pathways as well as the expression and functionality of IFN-stimulated gene (ISG) products and other key sites of host defence mechanisms such as induction of apoptosis within the infected cell. Therefore, HCV might successfully adapt to the host environment and evade the host antiviral response, leading to persistent viral replication and spread.⁸ The key innate immune evasion strategies used by HCV to establish persistent infection within the liver as well as how host genotype influences the outcome of HCV infection, was reviewed by Horner. ⁹ The key viral factor involved in the HCV immune evasion programme is the viral NS3/4A protease, which cleaves mitochondrial antiviral signalling protein (MAVS), the RIG-I signalling adaptor protein and Tolllike receptor-domain containing adapter-inducing IFN-β (TRIF), the TLR3 signal transducer. Additionally, there is an increasing role for microRNAs in mediating HCV innate immune evasion strategies, including m iR-122¹⁰ and miR-21.¹¹

HIV

Chronic HIV infection is linked with a variety of immune evasion strategies beyond the basic integration of the HIV-1 genome into target cells. These include an overall ability to maintain a latent infection and to avoid antibody responses by varying or shielding immunodominant epitopes within the viral envelope glycoprotein. Latent HIV resides in memory T cells or myeloid cells (HIV reservoirs), persisting indefinitely in patients even on potent antiretroviral therapy. The molecular mechanisms of HIV latency are complex and include the absence in resting $CD4⁺$ T cells, inhibition of nuclear forms of key host transcriptional factors (e.g. nuclear factor- κ B and nuclear factor of activated T cells), the presence of transcriptional repressors, blocking of mRNA splicing, insufficient Tat activity and associated host factors that promote efficient transcriptional elongation, and transcriptional interference, as reviewed previously.^{12–14}

The HIV envelope glycoprotein that drives HIV entry is highly variable. Its plasticity allows HIV to escape the immune system, and its variability is associated with HIV tropism, fitness and replicative capacity. Entry inhibitors represent a new generation of antiviral for the treatment of HIV infection. Several compounds which block the attachment of HIV gp120 to either the $CD4^+$ T-cell receptor or the CCR5/CXCR4 co-receptors are currently in clinical development. Changes within a 10-amino-acid segment encompassing residues 36–45 within the HR1 region of gp41, however, may cause resistance to the entry inhibitor, enfuvirtide, while multiple changes in different gp120 domains (V1, V2, V3, C2 and C4) are associated with loss of susceptibility to other entry inhibitors.¹⁵ Resistance to antibodies by point mutation on the V2/V3 loop and N-linked carbohydrate glycans on gp120 has been indicated as hindering antibody neutralization.¹⁶ Recently, one study found that $CD4^+$ T memory stem cells harbour high per-cell levels of HIV-1 DNA and make increasing contributions to the total viral $CD4^+$ Tcell reservoir over time. Hence, HIV-1 may exploit the stem cell characteristics of cellular immune memory to promote long-term viral persistence.¹⁷

Hepatitis B virus

During active replication, HBV produces enormous viral loads in the blood and a massive surplus of subviral surface antigen particles in the serum of infected patients without killing hepatocytes. Together with the use of a reverse transcriptase during replication, it provides an enormous genetic flexibility for the selection of viral mutants upon selective pressure.¹⁸ Escape mutations in the major hydrophilic region of hepatitis B surface antigen (HBsAg) are reported widely, and these mutations lead to diagnostic problems, emergence of vaccine-escape mutants, and hepatitis B immunoglobulin prophylaxis failure.¹⁹ For example, mutations in the preS gene can result in undetectable HBsAg even when HBV is replicating. Surface gene mutations lead to decreased binding affinity to anti-HBs, which are associated with a vaccine escape mutant that is occasionally found in clinical cases. Mutations in the basal core promoter are associated with increased HBV replication and high incidence of progressive liver diseases. HBx protein can disrupt RIG-I-mediated IFN- β induction by down-regulating MAVS²⁰ and acts as a de-ubiquitinating enzyme to de-ubiquitinate RIG-I and other molecules. 21 One research study suggests that HBV polymerase (Pol) blocks IFN regulatory factor (IRF) signalling, indicating that HBV Pol may be the viral molecule that effectively counteracts host innate immune response in the early phase of the infection.²² It appears that HBV has evolved numerous strategies to counteract defence mechanisms and maintain persistence by actively manipulating the host innate immune response. 23,24

Host defence mechanisms for persistent infection

For many viruses, it has been shown that innate immune responses, most notably induction of type I and III IFNs, are the first line of defence limiting viral replication and spread, so contributing to the outcome of an infection. Given the importance of this defence, viruses have evolved numerous counteracting strategies including the blockade of IFN induction, interference with signalling triggered by IFNs, inhibition of the action of ISGs, or disturbance of the action of IFN-induced antiviral proteins.^{25,26}

Hepatitis C virus and interferon

Interferon- α is an essential component of innate antiviral immunity and of treatment regimens for chronic HCV infection. Resistance to IFN might be important for HCV persistence and failure of IFN-based therapies. Several in vitro escape experiments led to the identification of HCV envelope mutations resulting in increased viral fitness and conferring IFN- α resistance.²⁷ Hepatitis C virus can shut down the host IFN responses by using the viral NS3/4A protease to cleave MAVS/virus-induced signaling adapter²⁸ and TRIF, 29 two key adaptor molecules essential for interferon signalling activation. Recently, a study showed that NS4B can block the interaction of stimulator of interferon genes and Tank-binding kinase 1 to evade host innate immunity. 30 A strong association with viral clearance was found for a single-nucleotide polymorphism (SNP) in the promoter/enhancer region of the IL28B gene (encoding IFN-k3). The strongest association was found for SNP rs12979860, which is located about 3 kb upstream of the IL28B coding region.

Patients with a CC genotype at this SNP were more than twice as likely to achieve a sustained virological response as patients with a CT or TT genotype. 31 Both the success rate of treatment and the rate of spontaneous virus clearance were shown to increase profoundly in patients with the C/C genotype. $31-34$ Recently, researchers discovered a new transiently induced region upstream of IFNL3 (IL28B) on chromosome 19q13.13 that harbours a dinucleotide variant ss469415590 (TT or ΔG) which is in high linkage disequilibrium with rs12979860, a genetic marker strongly associated with HCV clearance. Compared with rs12979860, ss469415590 is more strongly associated with HCV clearance in individuals of African ancestry, although it provides comparable information in Europeans and Asians.³⁵ Moreover, the IFNL4 ss469415590 variant was identified as a better predictor than IFNL3 (IL28B) rs12979860 of pegylated IFN-a/ribavirin therapy failure in HCV/HIV-1 co-infected patients.³⁶

HIV/hepatitis B virus and interferon

HIV persistence is linked with the ability of the virus to dysregulate and evade the innate immune response. The IRFs are transcription factors that play major roles in innate immunity. Rustagi and Gale reviewed the interplay between HIV-1 and IRF3.³⁷ It was reported that Vpu plays a role in antagonism of IRF3 function.³⁸ However, Vpu was later reported to inhibit the activation of the IFN- β promoter by blocking nuclear factor- κ B signalling instead of IRF3. 39 A frequent functional TLR7 polymorphism, TLR7 Gln11Leu, resulting in significantly less IFN- α production, has been associated with higher viral loads and accelerated disease progression to advanced immune suppression in HIV patients and may also be associated with increased HIV susceptibility.⁴⁰ The expression of HBV polymerase (Pol) in human hepatic cell lines was also found to inhibit the induction of IFN-stimulated genes and to result in the weakened antiviral activity of IFN-a. This provided a possible molecular mechanism by which HBV resists IFN therapy and maintains its persistence. 41 Kimkong et al.⁴² showed that the A allele of $1823G/A$ SNP within the IFNA1 gene was significantly associated with an increased risk of chronic HBV infection compared with healthy individuals and the self-limited HBV group. Furthermore, HBV induces the expression of interleukin-8 (IL-8), which in turn reduces HBV sensitivity to IFN- α .⁴³ All HBV protein components can reportedly impair type I IFN transcription in different ways. The molecular mechanism underlying the HBV-induced suppression of type I IFN-mediated antiviral immunity was reviewed by Guo. ⁴⁴

Receptors and ISGs

Numerous reviews have detailed the host immune system control of viral infection.^{45–47} The innate immune system constitutes the first line of defence against viruses, initiating an antiviral response. Viruses are recognized by this system primarily through detection of their nucleic acids, either their packaged genome or viral replication intermediates within the infected cell.⁴⁸ Three classes of receptors, designated RIG-I-like receptors, TLRs and nucleotide oligomerization domain-like receptors, sense viral components such as double-stranded RNA, single-stranded RNA and DNA.⁴⁹ These types of recognition induce the transcription of proinflammatory cytokines and type I IFNs, inducing the expression of hundreds of ISGs, which engage in counteracting virus replication and spread. However, viruses have evolved a fine-tuned mechanism to evade detection by the immune system or to interfere with the resulting signalling pathways as described previously.50–⁵³

APOBEC3

The apolipoprotein B mRNA-editing enzyme catalytic polypeptide 3 (APOBEC3) subgroup plays an important role in the innate immune system, acting in host defence against exogenous viruses and endogenous retro-elements. The role of APOBEC3 proteins in the inhibition of viral infection was first described for HIV-1. However, in the past few years many studies have also shown evidence of APOBEC3 action on other viruses associated with human diseases, including HCV, HBV, human T-lymphotropic virus, human papillomavirus, herpes simplex virus-1, and Epstein–Barr virus. APOBEC3 inhibits these viruses through a series of editing-dependent and -independent mechanisms. However, many viruses have evolved mechanisms to counteract APOBEC effects.⁵⁴

Autophagy

Autophagy is an evolutionarily conserved intracellular process by which bulk cytoplasm is enveloped inside a double-membraned vesicle and shuttled to lysosomes for degradation. Autophagy is essential for tissue homeostasis and development, and defective autophagy is associated with a number of diseases. With respect to their role in antiviral responses, the autophagy proteins function in targeting viral components or virions for lysosomal degradation in a process termed xenophagy, and they also play a role in the initiation of innate and adaptive immune system responses to viral infections. Consistent with this antiviral role of host autophagy, some viruses encode virulence factors that interact with the host autophagy machinery and block the execution of the process. In contrast, other viruses appear to use components of the autophagic machinery to foster their own intracellular growth or non-lytic cellular egress.⁵⁵

Autophagy has virus-specific roles relating to viral replication, host innate and adaptive immune responses, virus-induced cell death programmes, and viral pathogenesis.⁵⁶ Autophagy has been proposed as a protective mechanism against viral infection by degrading the pathogens within autolysosomes. This is strengthened by the fact that several proteins involved in IFN signalling pathways are linked to autophagy regulation. Again, several viruses have evolved strategies to divert IFN-mediated pathways and autophagy to their own benefit.⁵⁷ A recent study showed that HCV perturbs mitochondrial dynamics by promoting mitochondrial fission followed by mitophagy, which attenuates HCV-induced apoptosis and contributes to persistent HCV infection.⁵⁸

Interactions of immune cells

Dendritic cells

Chronic HBV or HCV infection is the result of an inadequate immune response towards the virus. Dendritic cells (DC) of patients with chronic HBV or HCV are impaired in their maturation and function, resulting in more tolerogenic rather than immunogenic responses, which may contribute to viral persistence. However, the experimental findings are controversial, some researchers finding no

quantitative, phenotypic, or functional impairment of myeloid DC (mDC) or plasmacytoid DC (pDC) in chronic HBV or HCV.

It has been demonstrated that both HBV particles and purified HBsAg have an immune modulatory capacity and may directly contribute to the dysfunction of mDC in patients with chronic HBV.⁵⁹ HBsAg inhibited the production of IFN-a by pDC through the induction of monocytes that secrete TNF- α and IL-10, and through the down-regulation of TLR9 expression on $pDC₆₀$ The direct immune regulatory effect of HBV and circulating HBsAg particles on the function of DC can be considered as part of the mechanism by which HBV escapes immunity.

Likewise, HCV inhibits cell surface expression of HLA-DR, prevents DC maturation, and induces IL-10 production. HCV can have direct and/or indirect inhibitory effects on antigen-presenting cells, resulting in reduction of antigen-specific T-cell activation.⁶¹ Studies have revealed that different HCV proteins used distinct mechanisms to down-regulate DC functions. Individual HCV proteins (Core, NS3, NS4, NS5) as well as fused polyprotein (Core-NS3-NS4) were found to impair functions of both immature DC and mature DC by regulating the expression of co-stimulatory and antigen presentation molecules, strikingly reducing IL-12 secretion, inducing the expression of Fas ligand to mediate apoptosis, interfering with allostimulatory capacity, inhibiting TLR signalling, and inhibiting nuclear translocation of nuclear factor- κ B in DC.⁶² In addition, DC inhibition is connected to exhaustion of CD8⁺ T-cell functionality, including production of IFN- γ , IL-2, TNF- α and CD107a mobilization during chronic HCV infection.⁶³ Myeloid DC from patients with chronic HCV infection expressed up-regulated levels of two inhibitory ligands, Fas ligand and the ligand 2 of programmed death-1 (PD-L2), compared with mDC from healthy people. The mDC of HCV patients had cytotoxic effects on autologous patient T cells and allogeneic healthy T cells. These results indicate that the cytotoxic activity of mDC is up-regulated to kill T cells during chronic HCV infection, which represents a novel mechanism of HCV immune evasion.⁶⁴ All of these effects may account for or contribute to the low overall level of immunogenicity of HCV observed in chronically HCV-infected patients.

Dendritic cells are key regulators of the host response to HIV-1 infection. They are among the first cells to encounter HIV-1 at mucosal sites where they are co-opted by HIV-1 to facilitate transmission. 65 The DC not only encounter the virus in the sexual mucosa, but, more importantly, they are also able to transfer the virus to its primary target cells, $CD4^+$ T cells, either within the mucosa or after migration to lymph nodes. This results in explosive viral replication that is much greater than that resulting from direct infection of T cells. 66 HIV-1

directly and indirectly modulates DC function to hinder the formation of effective antiviral immunity and fuel immune activation. HIV-1 evades innate immune sensing by mDC, resulting in suboptimal maturation and poor generation of antiviral adaptive responses and contributing to regulatory T (Treg) cell development.

Plasmacytoid DC interactions with HIV-1 are pleotropic, modulating immune responses on an axis between immunostimulatory and immunosuppressive. The pDC promote immune activation through an altered phenotype of persistent type I IFN secretion and weak antigen presentation capacity. Conversely, HIV-1 stimulates secretion of indoleamine 2,3 dioxygenase by pDC, resulting in Treg cell induction. 67 This not only serves to blunt anti-HIV immune responses but also may dampen chronic immune activation. Dysfunction of DC in chronic persistent infectious disease is multifactorial, implicating a combination of viral and non-viral factors. We presume that viral persistence is the result of the host's insufficient antiviral immune responses and viral evasion.

Natural killer cells

Natural killer cells are important antiviral effectors of innate immunity because of their contribution to virus elimination via direct killing of infected cells and cytokine (IFN- γ and TNF- α) production; their regulation depends on a finely tuned balance between inhibitory and activating receptors. Recent research highlights the fact that NK cells are also regulatory cells engaged in reciprocal interactions with DC, macrophages, T cells and endothelial cells. Hence, NK cells can limit or exacerbate immune responses.^{68,69}

In the last few years major progress has been made in better understanding the role of NK cells in HCV infection. This includes multiple pathways by which HCV impairs or limits NK cell activation. Many studies showed that chronic HCV infection is associated with dysfunctional or biased NK cell phenotypes. Compelling studies suggested that peripheral NK cells are activated during chronic HCV infection, most likely by IFN- α/β signalling. These activated peripheral NK cells display an increase in cytotoxicity with elevated expression of NKG2D, NKp46, and TNF-related apoptosis inducing ligand (TRAIL) and elevated activation of signal transducer and activator of transcription 1 (STAT1) along with a decrease in IFN- γ production.^{70,71} The elevated cytotoxicity level of NK cells may contribute to liver injury, whereas the decreased production of IFN- γ may facilitate the inability to clear HCV.⁷²

Previous studies showed that the cross-linking of the E2 protein of HCV with CD81 on NK cells inhibits their activation, cytokine production, cytotoxic granule release and proliferation. These results implicate HCV-E2-mediated inhibition of NK cells as an efficient HCV evasion strategy targeting the early antiviral activities of NK cells and allowing the virus to establish itself as a chronic infection. Although the data are still controversial, 73 studies suggested that genes encoding the inhibitory NK cell receptor KIR2DL3 and its human leucocyte antigen C group 1 (HLA-C1) ligand directly influence resolution of HCV infection.⁷⁴ The specific combination of KIR2DL3 and HLA-C1 was protective against chronic HCV infection. Natural killer cells contribute to anti-HCV defence in vivo in the earliest stages of infection, providing innate protection from HCV acquisition.⁷⁵ Genetic and clinical studies have suggested that NK cells not only play an important role in both spontaneous and IFN- α therapybased HCV clearance, but may also contribute to hepatocellular damage in viral hepatitis.^{76,77}

Like NK cells in HCV patients, NK cells in HBV patients enhance cytolytic activity and dysfunctional cytokine production, which may contribute to virus persistence78,79 and liver inflammation by TRAIL-mediated death of hepatocytes.^{80,81} Recent data have highlighted that NK cells are capable of exerting antiviral and immunoregulatory functions while also contributing to the pathogenesis of liver injury via death receptor pathways like a double-edged sword.^{82,83}

Studies have suggested that NK cells can contribute to the control of HIV-1 infection through the recognition of virus-infected cells by both activating and inhibitory killer immunoglobulin-like receptors (KIRs) like KIR3DS1 and KIR3DL1.84,85 The NK cells expressing KIR3DS1 showed strong, significant dose- and cell contact-dependent inhibition of HIV-1 replication in target cells expressing HLA-B Bw4-80I compared with NK cells that did not express KIR3DS1.⁸⁶ However, studies demonstrated that KIR+ NK cells can place immunological pressure on HIV-1 and that the virus can evade such NK cell-mediated immune pressure by selecting for sequence polymorphisms. Therefore, NK cells may play a role in contributing to viral evolution. 87 Several key NK cell receptors have been associated with HIV-1 disease progression and/ or transmission, implying that NK cells might contribute markedly to the control of HIV-1 infection⁸⁸ while HIV-1-encoded proteins can facilitate evasion from NK cell recognition.⁸⁹

Regulatory T cells

Surviving an infection requires the generation of an immune response that controls the invading pathogen while limiting collateral damage to self tissues that may result from an exuberant immune response. Various populations of regulatory cells, including CD4+ CD25+ Foxp3⁺ Treg cells, have been shown to play a central role in the establishment of these controlled immune responses. Treg cells are involved in the control of immune-tolerance by regulating immune homeostasis and limiting immune activation. Defects in Treg cell numbers

or function have been related to the development of human autoimmune diseases, while increases in Treg cell numbers or activity could limit anti-tumour immune responses.⁹⁰

In contrast to these two scenarios, a much more complex picture emerges for the role of Treg cells in infectious diseases. Treg-cell-mediated inhibition of antimicrobial immune responses could lead to ineffective clearance of the pathogen contributing to the chronicity of the infection. Conversely, Treg cells participate in terminating immune responses, so preventing exacerbated and potentially harmful immune activation and immunemediated injury. The balance between regulatory T cells and effector immune functions influences the outcome of host–virus coexistence and appears tuned to maintain immune homeostasis.⁹¹

HIV infection is associated with a progressive CD4⁺ T-cell lymphopenia, and defective HIV-specific CD8⁺ T-cell responses are known to play a key role in the control of viral replication. Persistent immune activation is a hallmark of HIV infection and is involved in disease progression independent of viral load. The consequences of Treg cell expansion, observed in HIV infection, could be either beneficial by suppressing generalized T-cell activation or detrimental by weakening HIV-specific responses and so contributing to viral persistence. The resulting balance between these contrasting outcomes might have critical implications in pathogenesis.^{92,93} Treg cell heterogeneity analysis in the context of HIV infection was reviewed by Simonetta and Bourgeois.⁹⁴ They discussed how the identification of naive and effector Treg cell subsets modulates understanding of Treg cell biology during HIV infection and the potential impact of HIV infection on mechanisms governing peripheral differentiation of adaptive Treg cells.

Treg cells also appear to play a role in the control of chronic viral hepatitis.⁹⁵ Patients with either chronic HBV or HCV infections have accumulated circulating Treg cells that express CD45RO, high intracellular cytotoxic T-lymphocyte antigen 4 (CTLA-4) and PD-1 and which have the capacity to inhibit proliferation of both CD4⁺ and $CD8⁺$ T cells as well as T helper type 1 (Th1) cytokine production. An increased number of circulating Treg lymphocytes has been shown in patients with chronic HBV infection when compared with those who have cleared HBV.⁹⁶ Similar findings have been found in chronic HCV infection when compared with healthy controls.⁹⁷ Of particular interest is that viral load and the number of Treg cells are closely related. Therefore, chronic HBV or HCV patients harbour an increased percentage of Treg cells in peripheral blood compared with controls. Treg cells have an immunosuppressive effect on HBV/HCV-specific T helper cells. The presence of Treg cells could contribute to an inadequate immune response against the virus, leading to chronic infection. Treg cells

generated in chronic viral infection may be part of a normal host immune response to infection with a natural evolution from pro-inflammatory responses to immune regulation, where neither are related directly to virus infection and where the benefit would be in preventing uncontrolled tissue destruction. Treg cells could, in that situation, be regarded as protective, with viral persistence and less liver damage, a by-product of that protection. However, viral induction of the Treg cell response favours persistent infection and survival of the organism. One could speculate that the hepatitis viruses skew the natural immune response towards regulation rather than inflammation as part of evolutionary development.⁹⁵ One consequence of the modulation of excessive immune responses and control of immunopathology by Treg cells is enhanced pathogen survival and possibly long-term viral persistence.

Hence, pathogen persistence may represent a compromise reached between the pathogen and the host immune response. Activation of Treg cells may contribute to the maintenance of immunity to chronic infections in which pathogen persistence is required for the development of memory. A balance between effector and regulatory mechanisms may determine the outcome of an infection, and this may, in some cases, be beneficial for both the host and the pathogen.^{98,99} In conclusion, Treg cells contribute to the viral persistence and maintain immune homeostasis.

Regulatory B cells

Emerging significant evidence indicates that B cells can actively modulate immune responses through mechanisms that do not directly involve the production of antibodies. B cells appear to have the capacity to both induce and suppress immune effector mechanisms, and they exert these functions both by contact-dependent interactions and through the secretion of cytokines. These B cells are functionally defined 'regulatory B cells', or Breg cells.¹⁰⁰ Breg cells may act earlier than Treg cells and may play an important role in autoimmune and allergic diseases. The Breg subtypes may be homologous to the Treg subtypes (Br1 cells expressing IL-10, Br3 cells expressing transforming growth factor- β , and B-Foxp3 cells), although the Br1 subtype seems to predominate. Nevertheless, differences with Treg cells may exist: Breg cell activation may chiefly involve the TLRs rather than the antigen receptor, and Breg cells act earlier, facilitating the recruitment of Treg cells and then disappearing once the Treg cells become operational.^{101,102} In autoimmunity, tumour immunology, and some viral infections, Breg cells modulate T-cell function via IL-10 production. Studies indicate that Breg cells contribute to HIV infection-associated immune dysfunction by T-cell impairment via IL-10 and possibly PD-L1 expression.¹⁰³

Interactions of cytokines and regulatory molecules

Programmed death-1

Viral infection may modulate many molecules to impair specific cytotoxic T lymphocyte reactivity through induction of either a virus-associated tolerogenic-like state or apoptosis. Virus-specific CD8⁺ T-cell exhaustion may represent a mechanism of viral persistence. One key pathway is the inhibitory receptor programmed death-1 (PD-1) binding to one of its ligands, PD-L1 or PD-L2, transmitting a negative signal to the T cells expressing PD-1, reducing cytokine production and proliferation, and regulating the balance between T-cell activation, tolerance and immunopathology.¹⁰⁴ PD-1 has been reported to be up-regulated in exhausted CD8⁺ T cells during chronic viral infections, such as HBV, HCV and HIV. Expression of PD-1, which is driven by viral replication and associated with T-cell dysfunction, was also up-regulated on virus-specific $CD4^+$ T cells.¹⁰⁵ Expression of PD-1/PD-L1 was also up-regulated in monocytes/macrophages in HCV-infected subjects compared with HCV-resolved or healthy subjects. Up-regulation of PD-1 was inversely associated with the degree of IL-12 inhibition in HCV infection.¹⁰⁶ Our research data suggested that HCV core/gC1qR engagement on monocytes/ macrophages triggers the expression of PD-1 and suppressor of cytokine signalling 1 (SOCS-1), which can associate to deliver negative signalling to TLR-mediated pathways controlling expression of IL-12, a key cytokine linking innate and adaptive immunity.¹⁰⁷ The inhibitory immunoregulatory receptor, CTLA-4, was also up-regulated in HIV-specific $CD4^+$ T cells but not $CD8^+$ T cells in all categories of HIV-infected subjects.108

Blocking PD-1/PD-L1 interaction with an anti-PD-L1 antibody improved the capacity of expansion of virusspecific CD4⁺ CD8⁺ T cells. However, highly PD-1-positive intrahepatic CD8⁺ T cells were more phenotypically exhausted with increased CTLA-4 and reduced CD28 and CD127 expression, suggesting that active antigen-specific stimulation in the liver induces a profound functional exhaustion not reversible by PD-1/PD-L blockade alone.¹⁰⁹ A PD-1⁺ CD127⁻ phenotype associated with anergic features and apoptosis in chronic HCV patients has been identified.¹¹⁰ In the case of $CD4^+$ T cells, blockade of PD-L1/2, IL-10 and transforming growth factor- β_1 increased expansion of $CD4^+$ T cells in patients with chronic HCV, restoring HCV-specific production of IL-2, IFN- γ and TNF- α .¹¹¹

Likewise, intrahepatic HBV-specific CD8⁺ T cells expressed higher levels of PD-1 and lower levels of CD127 than their peripheral counterparts. Blockade of PD-1/PD-L1 interaction increased CD8⁺ T-cell proliferation, IFN- β and IL-2 production by circulating intrahepatic lymphocytes, even though anti-PD-L1 had a

stronger effect on intrahepatic compared with peripheral T cells.¹¹² Anti-PD-L1 might be a good therapeutic candidate for chronic viral infection.

Tim-3

Notably, not all dysfunctional T cells express PD-1, nor are they all rescued by blockade of the PD-1/PD-1 ligand pathway. The expression of T-cell immunoglobulin and mucin domain-containing protein 3 (Tim-3) is increased on CD4⁺ and CD8⁺ T cells in chronic virus infections such as HBV, HCV and HIV. Tim-3 expression correlates with a dysfunctional and senescent phenotype (CD127^{low} CD57^{high}), a central rather than effector memory profile (CD45RA^{negative} CCR7^{high}), and reduced Th1/ Tc1 cytokine production.¹¹³ Up-regulation of Tim-3 and inhibition of IL-12 are also observed in monocytes/macrophages incubated with HCV-expressing hepatocytes, as well as in primary monocytes/macrophages or monocytic THP-1 cells incubated with HCV core protein. Tim-3 blockade reduces HCV core-mediated expression of the negative immunoregulators PD-1 and SOCS-1 and increases STAT-1 phosphorylation. Conversely, blocking PD-1 or silencing SOCS-1 gene expression also decreases Tim-3 expression and enhances IL-12 secretion and STAT-1 phosphorylation.¹¹⁴

The role of PD-1 and Tim-3 expression in T-cell exhaustion during HCV/HIV co-infection has been investigated. HCV-specific T cells in HCV/HIV co-infection show elevated frequencies of dual Tim-3/PD-1 expression that correlate with liver disease progression.¹¹⁵ Several markers of immune exhaustion such as PD-1, Lymphocyte Activation Gene-3, Tim-3 and CTLA-4, which are also negative regulators of immune activation, are preferentially up-regulated on T cells during HIV infection, 116 and in vitro blockade of PD-1 and Tim-3 restores specific T-cell responses.

In addition, the Tim-3 pathway appears to control Treg cell and effector T (Teff) cell balance by altering cell proliferation and apoptosis during HCV infection. 117 Blockade of Tim-3 on $CD4^+$ CD25⁺ T cells promoted expansion of Teff cells more substantially than Treg cells through improved STAT-5 signalling, so correcting the imbalance of $F\exp 3^{+}$ Treg $F\exp 3^{-}$ Teff cells that was induced by HCV infection. Beyond T cells, there was a significant increase of Tim-3 expression in peripheral blood mononuclear cells, circulating NK cells, and liver infiltrating lymphocytes from chronic HBV patients compared with healthy controls, which may in turn suppress NK cell functions in chronic HBV patients.¹¹⁸

NKG2A

With regard to NK cells, expression levels of NKG2A were up-regulated on NK cells from individuals with

chronic HCV. Treatment with pegylated IFN and ribavirin resulted in a down-regulation of NKG2A expression on CD56^{dim} NK cells whereas individuals with a sustained virological response had greater numbers of NKG2A-positive, KIR-negative NK cells than those without sustained virological response. 119 In addition, infection with HBV increases levels of the inhibitory receptor NKG2A on NK cells in humans and reduces their ability to clear HBV. Reagents designed to block the interaction between NKG2A and HLA-E in peripheral NK cells from patients with active chronic HBV increased their cytotoxicity in vitro.¹²⁰ Higher NKG2A expression levels in the cytotoxic NK subset were found in the later stages of HIV infection compared with patients at an early stage of infection. A reverse association between the percentage of NKG2A-positive cells in the cytotoxic NK subset and the CD4 cell count was observed in all HIV-1-infected groups.121 NKG2A seems to be a target of viral persistence and a modulator of NK immune activation.

KLRG1

We demonstrated that killer cell lectin-like receptor subfamily G member 1 (KLRG1), a transmembrane protein preferentially expressed on T cells, is highly expressed on CD56+ NK cells, which are significantly reduced in their numbers and functions in the peripheral blood of patients with chronic HCV infection compared with subjects without infection. Importantly, blockade of KLRG1 signalling significantly recovered the impaired IFN- γ production by NK cells from HCV-infected subjects.¹²² We also demonstrated that KLRG1 was over-expressed on CD4⁺ T cells from HCV-infected, HBV vaccine non-responders compared with HBV vaccine responders. These results may partly explain why HBV vaccine responses in HCVinfected individuals are often blunted when compared with uninfected populations.¹²³

Interleukin-17

Interleukin-17A (also called IL-17) is the prototypic cytokine of the IL-17 family, which includes six members: IL-17A, B, C, D, E and F. IL-17A and IL-17F have similar functions: they induce the production of pro-inflammatory cytokines, chemokines and metalloproteinases from various tissues and cell types. Researchers found that peripheral blood Th17 cell levels in patients with chronic hepatitis B were significantly higher than those in the healthy controls associated with elevated serum levels of IL-17A.124,125 Increased IL-17-secreting cells in the liver and peripheral Th17 cells among patients with chronic hepatitis B are associated with greater fibrosis and higher degrees of inflammation compared with healthy controls.¹²⁶ In vitro, IL-17 together with IL-17-activated monocytes were able to promote the activation of stellate

cells, which in turn exacerbates liver fibrosis and the inflammatory response.¹²⁷ Interleukin-17-mediating liver neutrophil recruitment via induction of IL-8 may be one potential mechanism of liver injury in patients with chronic HBV.¹²⁸

Both Th17 frequency and plasma IL-17A levels inversely correlated with viral load in chronic HBV infected patients.¹²⁹ Wang et al. demonstrated that IL-17A effectively suppressed HBV replication in a non-cytopathic manner, and the over-expression of myxovirus resistance A and oligoadenylatesynthetase mRNA was involved in the suppression of HBV replication by IL-17A.¹³⁰ All these findings suggest that IL-17 participates in the liver injury process and viral clearance after HBV infection, which seems paradoxical. Other researchers found that the frequency of Th17 cells display no correlation with serum HBV DNA loads or alanine aminotransferase levels, although the frequency of Th17 cells in peripheral blood was significantly higher in paediatric patients with CHB compared with healthy controls.¹³¹

Serum IL-17 levels are higher in chronic HCV-infected patients, and HCV-specific Th17 cells correlated with liver damage but not HCV viral replication.¹³² However, novel intriguing data indicate that a Th17 boost could be associated with spontaneous HCV clearance.¹³³ It is possible that Th17 could play a dual role (both beneficial and harmful) and that an imbalance of regulating factors (chemokines, transcription factors, receptor expression) rather than the lymphocyte itself could tip the Th17 immune response one way or the other. The role of Th17 cells in host anti-HCV defence is beginning to emerge, and one has to focus upon its potential beneficial aspects and not only on its destructive potential.¹³⁴

During the course of HIV infection, Th17 cells are lost very early and their loss has been shown to correlate with bacterial translocation.¹³⁵ Interleukin-17 production was significantly lower in patients with detectable plasma viraemia when compared with successfully treated HIVinfected patients and healthy controls.¹³⁶ A significant negative correlation between virus-specific Th17 cells and HIV-1 plasma viral load (pVL) indicates a gradual loss of Th17 cells with HIV-1 disease progression.¹³⁷ The IL-17 production in HIV-infected patients could be recovered through a sustained suppression of viral replication in the peripheral blood through combined anti-retroviral therapy. When Th17 differentiation cytokines were given along with lamivudine to the cultures with established HIV infection, a Th17 response was fully restored and virus replication was suppressed.^{136,138,139} All these findings indicate the potential use of immunotherapeutic modalities to supplement anti-retroviral drugs for restoring Th17 response in chronically infected patients.

In summary, diverse regulatory mechanisms cooperate to maintain immune homeostasis, and a breakdown in these pathways may facilitate viral persistence. Use of immunotherapeutic modalities to supplement anti-retroviral drugs may contribute to immune homeostasis and prevent viral damage to the host in chronically infected patients.

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Disclosures

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