Interferon- λ in the Context of Viral Infections: Production, Response and Therapeutic Implications

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Key Words

Epithelium · Hepatitis C virus · Host defense · Interferon · IFNAR · IFNLR1 · IL-28 · IL-29 · RNA virus · Virology

Abstract

Interferon (IFN)- λ forms the type III IFN family. Although they signal through distinct receptors, type I (IFN- α/β) and type III IFNs elicit remarkably similar responses in cells. However, in vivo, type III and type I IFN responses are not fully redundant as their respective contribution to the antiviral defense highly depends on virus species. IFN- λ is much more potent than IFN- α/β at controlling rotavirus infection. In contrast, clearance of several other viruses, such as influenza virus, mostly depends on IFN- α/β . The IFN- λ receptor was reported to be preferentially expressed on epithelial cells. Cells responsible for IFN- λ production are still poorly characterized but seem to overlap only partly IFN- α/β -producing cells. Accumulating data suggest that epithelial cells are also important IFN- λ producers. Thus, IFN- λ may primarily act as a protection of mucosal entities, such as the lung, skin or digestive tract. Type I and type III IFN signal transduction pathways largely overlap, and cross talk between these IFN systems occurs. Finally, this review addresses the potential benefit of IFN- λ use for therapeutic purposes and summarizes recent results of genome-wide association studies that identified polymorphisms in the region of the IFN- λ 3 gene impacting on the outcome of treatments against hepatitis C virus infection.

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Introduction

Ten years ago, interferon (IFN)- λ was discovered by two independent groups [1, 2]. IFN- λ forms the type III IFN family, which is composed of three members, IFN- λ 1, IFN- λ 2 and IFN- λ 3, also named IL-29, IL-28A and IL-28B, respectively [1, 2]. Recently, a fourth IFN- λ subtype, named IFN- λ 4, has been described in humans [3]. It is expressed in a small fraction of the human population as a consequence of a frameshift occurring in a coding sequence, upstream of the *IFNL3* gene. This IFN only bears 29% identity with IFN- λ 3 but acts through the same receptor and displays typical antiviral activity [4].

Genes encoding IFN- λ map on chromosome 19q13 in humans and 7A3 in mice [1, 2, 5]. In mice, IFN- λ 1 is a pseudogene [5]. Unlike most type I IFN genes, which are intronless, type III IFNs have several exons, 5 for IFN- λ 1 and 6 for IFN- λ 2 and IFN- λ 3 [2]. In humans, IFN- λ 4 was shown to be N-glycosylated and IFN- λ 1 contains a potential N-glycosylation site [4, 6]. In mice, both IFN- λ 2 and IFN- λ 3 are N-glycosylated [5].

Type III IFNs signal through a heterodimeric cell surface receptor composed of two chains: IFNLR1, which is specific to IFN- λ , and IL10RB, which is shared by other IL-10-related cytokines [1, 2, 7]. Type I IFNs use a different receptor (IFNAR), which is composed of subunits IFNAR1 and IFNAR2c. However, upon binding their cognate receptors, type I and type III IFNs trigger the

same Jak-STAT signal transduction pathway [5, 8, 9] (fig. 1), leading to the upregulation of many genes, called IFN-stimulated genes (ISGs) [reviewed in 10]. More than 300 ISGs have been identified. Their products act to control viral infection, to modulate immune responses or to control mitosis. In addition, some ISGs encode factors that participate in the IFN signal transduction pathway and, therefore, control IFN expression and response through positive or negative feedback loops (fig. 1).

When type I and type III IFN receptors are expressed by a single cell line, triggering either receptor complex leads to the upregulation of the same set of ISGs [11]. Therefore, a major question that arose after the discovery of IFN- λ was whether type I and type III IFN responses were redundant *in vivo*.

Antiviral Activity of IFN- λ in Mice

One of the first evidence-based studies reporting IFN- λ antiviral activity *in vivo* was published by Ank et al. [8], who observed that intravaginal treatment of female mice with IFN- λ prior to infection with herpes simplex virus (HSV)-2 prevented virus replication in the vaginal mucosa. Next, analysis of IFN- λ antiviral activity *in vivo* largely benefited from the development of IFN- λ receptor-deficient mice (IFNLR1^{0/0}) [12]. By comparing viral infection in wild-type, IFNLR1^{0/0}, IFNAR^{0/0} and double-knockout mice, it was shown that the contribution of IFN- λ to the control of viral infection greatly varied according to the virus [13]. Table 1 and figure 2 recapitulate currently available data.

In spite of the efficacy of IFN- λ treatment observed against HSV-2, IFNLR1 deficiency did not substantially affect HSV-2 replication. In contrast, IFNAR1 deficiency strongly increased HSV-2 replication [12]. Thus, efficacy of IFN- λ against HSV-2 infection *in vivo* was modest compared to that of type I IFN. Additional studies showed no protective effect of IFN- λ against infection with encephalomyocarditis virus, lymphocytic choriomeningitis virus or vesicular stomatitis virus [8, 12]. Similarly, no antiviral activity of IFN- λ could be detected in mice against hepatotropic viruses such as Rift Valley fever virus, Lassa fever virus or a mutant strain of Thogoto virus (THOV) lacking the IFN-antagonist ML protein (Δ ML) [14, 15]. The lack of IFN- λ antiviral activity against these viruses was not due to a lack of IFN- λ expression in infected organs since IFN- λ was transcriptionally active in the liver of mice infected with THOV- Δ ML [14]. Yet, in spite of IFN- λ expression, THOV and Rift Valley fever

virus failed to induce Mx1 expression in infected livers of Mx^{+/+} mice lacking functional type I IFN receptors (IFNAR^{0/0}) [14].

IFN- λ was found to have a modest but detectable antiviral activity *in vivo* against influenza virus, human metapneumovirus and severe acute respiratory syndrome coronavirus. For these viruses, mice lacking both receptors turned out to be significantly more susceptible than IFNLR1^{0/0} mice [14–16]. Additional studies carried out with influenza virus showed variable extents of protection mediated by IFN- λ : two studies reported that IFN- λ was of moderate importance against influenza virus (strains SC35M, PR8- Δ NS1, A/HH/05/2009 and B/Lee/40) [14, 15] and one study observed a major role for IFN- λ (strain A/HK-X31) [16]. Such differences in influenza virus control might relate to variations in the ability of the different virus strains to spread systemically from the respiratory epithelium [16].

Contribution of IFN- λ was much clearer in the case of respiratory syncytial virus [15]. Indeed, control of respiratory syncytial virus infection depended equally on the presence of type I and type III IFN receptors. Interestingly, IFN- λ was shown to play a major, nonredundant, role in the protection against rotavirus infection [17]. IL-28R α ^{0/0}, and double-knockout mice were highly susceptible to oral rotavirus infection, while IFNAR^{0/0} and wild-type mice were resistant. Moreover, administration of IFN- λ allowed the control of rotavirus infection in mice, while administration of IFN- α did not. IFN- λ thus appears to contribute much more than type I IFNs to the protection against rotavirus [17].

Cellular Response to IFN- λ and Receptor Distribution

The receptor chain that is specific for IFN- λ was first identified in humans [1, 2, 7]. It has been discovered simultaneously by three different teams and called IL-28R α [2], LICR2 [7] or CRF2-12 [1]. It is now most commonly referred to as IFNLR1. By associating with the β chain of the IL-10 receptor (IL10RB), it forms the heterodimeric receptor necessary for IFN- λ signaling. The murine receptor has also been characterized [5]. Mouse IFNLR1 shares about 67% similarity with the human receptor and is encoded by a gene present on mouse chromosome 4D3. Interestingly, unlike type I IFNs, mouse and human IFN- λ are not species specific and can bind the receptors of both species [5].

Fig. 1. Type I and type III IFN signal transduction pathways. Viral nucleic acids are recognized by transmembrane TLRs, cytoplasmic DNA sensors and RNA helicases, leading to the activation of kinases. These kinases promote the activation of the NF- κ B, IRF3 and IRF7 transcription factors and their subsequent translocation to the nucleus where they stimulate IFN gene transcription. IFN- λ 1 and IFN- β gene expression largely depends on IRF3 and NF- κ B. Expression of IFN- λ 2 and IFN- λ 3, like that of IFN- α , depends more on IRF7 availability. Type I IFNs use a dimeric receptor composed of IFNAR1 and IFNAR2c. Type III IFNs signal through a different receptor, which is composed of IFNLR1 and IL10RB. Upon binding to their cognate receptors, type I and type III IFNs induce the same Jak/STAT pathway: the transphosphorylation and

activation of receptor-associated Jak1 and Tyk2 leads to the phosphorylation of STAT1 and STAT2 transcription factors. Phosphorylated forms of STAT1 and STAT2 further associate with IRF9 to form a heterotrimeric ISGF3 complex. ISGF3 then translocates to the nucleus where it binds to sequences of IFN-stimulated response elements present in the promoter of ISGs to upregulate their transcription. Some ISG products participate themselves in the signaling pathways leading to IFN production and IFN responses thus creating positive (and negative) feedback loops. Given the similarity of type I and type III IFN pathways, IFN- λ is expected to influence both the production of and response to IFN- α/β , and vice versa. cGAS = Cyclic GMP-AMP synthase.

Fig. 2. Relative contributions of type I and type III IFNs in antiviral protection: cartoon illustrating the relative weight of IFN-λ versus IFN-α/β in the control of viral infection. These data were deduced by comparing viral loads in mice deficient for either receptor. HMPV = Human metapneumovirus; RSV = respiratory syncytial virus; SARS-CoV = severe acute respiratory syndrome coronavirus.

Table 1. IFN-λ efficacy against viral infections

Effect	Virus	Route of infection	Organ (tissue) analyzed	Reference No.
No effect observed	Encephalomyocarditis virus	i.p.	Heart, brain	8, 12
	Lymphocytic choriomeningitis virus, <i>Armstrong strain</i>	i.v.	Spleen	8, 12
	Vesicular stomatitis virus, <i>Indiana strain</i>	i.v.	Spleen	12
	THOV	i.p.	Liver	14
	Rift Valley fever virus	i.p.	Liver	14, 15
	Lassa fever virus	i.n.	Liver	15
Moderate	HSV-2	i.vag.	Vaginal mucosa	8, 12
	Influenza	i.n.	Lungs	
	Influenza A SC35M (H7N7)			14
	Influenza A SC35M-ΔNS1			14
	Influenza A PR8-ΔNS1 (H1N1)			14
	Influenza A/HH/05/2009 (H1N1)			15
	Influenza B/Lee/40			15
	Influenza A/HK-X31 (H3N2)			16
	Human metapneumovirus	i.n.	Lungs	15
	SARS-CoV	i.n.	Lungs, intestine	15
Equal to type I IFNs	Respiratory syncytial virus	i.n.	Lungs	15
Major	Rotavirus	p.o.	Intestine	17

i.n. = Intranasal; i.p. = intraperitoneal; i.v. = intravenous; i.vag. = intravaginal; p.o. = peroral; SARS-CoV = severe acute respiratory syndrome coronavirus.

Epithelial Specificity of the IFN-λ Response

Unlike the type I IFN receptor, which is ubiquitously expressed, the type III IFN receptor – especially the IFNLR1 chain – displays a much more restricted cellular distribution [1, 2, 5, 18].

Several studies examined the responsiveness of human and mouse cell lines and primary cells to IFN-λ (table 2). Unsurprisingly, IFN-λ responses correlated with IL-28Ra expression [11]. Fibroblasts, splenocytes, bone-marrow-derived macrophages and endothelial cells did not re-

spond to IFN- λ , although they responded to IFN- α . In contrast, mouse melanoma cells and keratinocytes responded to IFN- λ [5]. Further studies demonstrated that epithelial cells are the primary targets of type III IFNs. Analysis of IFNLR1 expression in isolated mouse cells revealed that keratinocytes and vaginal epithelial cells, but not fibroblasts, were responsive to IFN- λ [12]. The first in vivo study of IFNLR1 expression in mice was performed in our laboratory [19]. In this study, the response to circulating IFN- λ was evaluated in vivo in various mouse organs after plasmid-mediated expression of IFN- λ from tibialis muscle cells. The tissues that showed the highest IFN- λ responsiveness – stomach, intestine, skin and lung – were epithelium-rich organs. Moreover, the response to IFN- λ paralleled IFNLR1 expression in analyzed tissues. In this study, the IFN- λ response was analyzed at the cellular level in kidney and brain sections using immunohistofluorescent detection of Mx1, a specific marker of the IFN response. IFN- λ -responsive cells strikingly differed from IFN- α -responsive cells. After IFN- λ expression, Mx1-positive cells almost exclusively corresponded to epithelial cells. These included cells forming the convoluted tubules in the kidney and the epithelial cells of the choroid plexus in the brain. In contrast to IFN- λ , circulating IFN- α induced a widespread response, which was pronounced in endothelial cells and included most cell types. The epithelial cell specificity of IFN- λ responses was subsequently observed in many other organs, such as the lung, intestine, stomach, skin or liver [15, 17, 19, 20] (fig. 3). In humans, constitutive expression of IFNLR1 was also observed in the upper epidermis [21].

The epithelial specificity of the IFN- λ response fits with the observed antiviral activity of this IFN in mouse infection models. The high impact of IFN- λ observed on rotavirus infection is in good agreement with the tropism exhibited by this virus toward intestinal epithelial cells.

It is noteworthy that the lack of antiviral IFN- λ activity observed for some viruses may stem from the use of intraperitoneal inoculation (table 1), which is expected to bypass the barrier made by the epithelial IFN- λ response.

Blood Immune Cell Responsiveness to IFN- λ

IFNLR1 is thus expressed in a very narrow range of cell types, mainly epithelial cells. However, some responses to IFN- λ were also reported in nonepithelial cells and notably in blood cells. Among these cells, conventional (cDCs) and plasmacytoid dendritic cells (pDCs) express IFNLR1 [22, 23] and IFN- λ was shown to alter their stimulatory properties [24, 25]. B cells and monocytes also express low but significant levels of IFNLR1 but fail to show de-

Table 2. Cells responsive to IFN- λ

Cells/organs	Reference No.
<i>Responsive cells (mouse)</i>	
Cell lines	
B16 melanoma cells	5
PAM212 keratinocyte-like cells	5
Primary cells	
Keratinocytes	5
Vaginal epithelial cells	12
pDCs	12
In vivo	
Kidney (epithelial cells)	19, 20
Epithelial cells of the choroid plexus	19
Intestine (epithelial cells)	17, 20
Airway cells	13–15, 20
Salivary glands	20
Cholangiocytes	30
<i>Responsive cells (human)</i>	
Cell lines	
HepG2, HuH7 (hepatocytes)	8, 18, 52, 62
HeLa	1
HaCaT (keratinocytes)	1
A594 (lung carcinoma cells)	1
Intestinal cell lines	1, 63
Primary cells	
Hepatocytes	18
Keratinocytes	21, 62, 64
Melanocytes	62
Natural killer cells	62
Airway epithelial cells	65
Nasal epithelial cells	66
Astrocytes	67
Neurons	67
pDCs	24
DCs	25
Monocytes	62
In vivo	
Liver	53, 54
<i>Nonresponsive cells (human and mouse)</i>	
Cell lines	
Fibroblasts	5
Primary cells	
Lymphocytes	12, 26
Monocytes, macrophages	5, 12, 62
Endothelial cells	62
Fibroblasts	12
Adipocytes	62
Splenocytes	5
In vivo	
Adipocytes	19
Endothelial cells	19
Hepatocytes (mouse)	30

Fig. 3. Epithelial specificity of the IFN- λ response in the intestine. Mx1 immunostaining in small-intestine sections of mice treated with IFN. **a** Response to circulating IFN- α in IFNLR1^{0/0} mice; Mx1 is mostly detected in lamina propria cells and little Mx1 expression is detected in epithelial cells. **b** Response to circulating IFN- λ in IFNAR^{0/0} mice; Mx1 expression is restricted to epithelial cells.

tectable responses. No expression was detected in natural killer and T cells [22].

IFN- λ Responses in the Liver

Hepatocytes are the most abundant cells in the liver and are targeted by several viruses. Because of their epithelial nature, hepatocytes are expected to respond to IFN- λ . Human hepatocyte cell lines and primary human hepatocytes indeed express the IFN- λ receptor and readily respond to IFN- λ [18, 26, 27]. Accordingly, IFN- λ was shown to restrict HCV replication in hepatoma cell lines. Clinical studies were conducted and IFN- λ entered phase 3 clinical trials as a candidate drug against HCV infection.

In contrast, in mice, response to IFN- λ appears to be very weak in the liver and IFNLR1 expression is hardly detectable in this organ [14, 19, 20]. Moreover, in mouse infection experiments, IFN- λ was not protective against hepatotropic viruses, such as Lassa fever virus, THOV or Rift Valley fever virus, despite documented production of IFN- λ in the liver of infected mice [14, 15]. Also, in transgenic mice harboring the genome of hepatitis B virus, IFN- λ was much less potent than IFN- β or IFN- γ at reducing viral replication [28].

The above data suggest that IFN- λ responses strongly differ between human and mouse hepatocytes. In a model of chimeric mice engrafted with human hepatocytes, the gene encoding IFNLR1 was more strongly expressed in human than in mouse hepatocytes [29]. Our own data

support these observations and show that human but not mouse hepatocytes respond to IFN- λ in chimeric mice treated with IFN- λ . In the mouse liver, IFN- λ responses were limited to cholangiocytes, the epithelial cells forming the bile ducts [30].

IFN- λ -Producing Cells

IFNs are produced after recognition of pathogen-associated molecular patterns by pattern recognition receptors. These receptors include the transmembrane Toll-like receptors (TLRs), cytoplasmic RNA helicases, such as RIG-I, MDA5 and LGP2, and cytoplasmic DNA sensors, such as the recently identified cyclic GMP-AMP synthase [31]. Pattern recognition receptors recognize byproducts of virus replication and trigger the synthesis of IFN. It was reported that type I and type III IFNs were induced by very similar signaling pathways [1, 2, 32, 33] (fig. 1). Indeed, IFN- λ 1 gene expression largely depends on IRF3 and NF- κ B, as that of IFN- β . IFN- λ 2 and IFN- λ 3 gene expression rather depends on IRF7, thus resembling that of IFN- α [33] (fig. 1). Yet, the range of IFN- λ -producing cells may not fully overlap that of type I IFN-producing cells. It was shown *in vivo* that the balance between type I and type III IFN gene transcription differed in the liver and the brain of infected mice [19].

After infection by HSV-2 or influenza virus, macrophages, which are good IFN- α/β producers, did not express IFN- λ [12, 34]. Thus, the cell types producing type I or type III IFNs partially overlap but are not identical. More recent studies have shown that the pathways leading to type I or type III IFN gene expression were not entirely identical, type III IFNs being more dependent on NF- κ B than type I IFNs [35, 36]. Expression of IFN- λ has been reported in DCs, respiratory epithelial cells, keratinocytes, hepatocytes, primary neuronal cells and a variety of cell lines [1, 12, 21, 34, 37–42].

IFN- λ Production by DCs

As in the case of type I IFNs, DCs and more particularly pDCs were shown to be important IFN- λ producers [22, 38, 43]. Upon viral infection with HSV-1, parapoxvirus or Sendai virus, pDCs produced large amounts of IFN- λ . However, it was reported that, in response to polyinosinic:polycytidylic acid [poly(I:C)], murine CD8 α ⁺ DCs, a subset of cDCs, were the major cell population producing IFN- λ [44]. Another study reported that human BDCA3⁺ cDCs, the counterpart of murine CD8 α ⁺ DCs, were the major cell population producing IFN- λ af-

ter treatment with poly(I:C) or after coculture with HCV-infected cells [23]. Interestingly, no release of IFN- α was detected from these cells.

In conclusion, both pDCs and cDCs produce IFN- λ . The fact that cells such as BDCA3+ cDCs produce IFN- λ but not IFN- α after stimulation by HCV-infected cells illustrates the subtle differences that exist in the pathways leading to type I or type III IFN production.

IFN- λ Production by Epithelial Cells

Epithelial cells are the main targets of IFN- λ activity. Interestingly, several studies suggest that epithelial cells may also act as IFN- λ -producing cells. IFN- λ was, for instance, the major IFN type induced by rhinovirus or influenza A virus infection of primary human airway epithelial cells [34, 45, 46]. In mice, IFN- λ production was observed in mouse tracheal epithelial cells in response to influenza A virus infection or poly(I:C) treatment [46].

IFN- λ production has also been reported in cultures of primary human keratinocytes in response to poly(I:C) treatment or vesicular stomatitis virus infection [21]. Interestingly, type I IFNs were only detected at very low levels in these cells.

Hepatocytes also produce IFN- λ . IFN- λ mRNA expression was induced in response to experimental HCV infection of human fetal liver cells or in patients with chronic HCV infection [42, 47]. More recently, in a model of chimeric mice transplanted with human hepatocytes, it was reported that IFN- λ was expressed and produced by human hepatocytes at a greater level than IFN- α or IFN- β in response to poly(I:C) treatment [29]. Interestingly, in the same study, human IFN- λ expression was more strongly induced in HepG2 cells than in HEK293T and MRC-5 cells (kidney and fibroblast cell lines, respectively), whereas the opposite pattern was observed for the expression of IFN- β . In mice, although the liver has proven to be weakly responsive to IFN- λ , this IFN was readily expressed in response to virus infection [14].

Finally, IFN- λ was also shown to be produced following bacterial infection with *Listeria monocytogenes* in vivo in the placenta of infected mice and in vitro by human intestinal cell lines, trophoblastic cells and HepG2 cells [48].

Interdependency of Type I and Type III IFNs

Several ISGs encode factors like RIG-like helicases, TLRs or IRF7, that participate in the signal transduction pathway leading to IFN production. Other ISGs encode proteins like STAT1, SOCS or Usp18, which positively or

negatively modulate IFN responses. Since type I and type III IFNs use very similar signal transduction pathways and upregulate the same group of ISGs, it is expected that cross talk exists between these two IFN systems (fig. 1).

Indeed, type III IFN expression was shown to be up-regulated by type I IFNs in conditions of virus infection. This was reported by different research groups for cDCs and macrophages, where treatment with IFN- α prior to virus infection increased IFN- λ expression [39, 40, 49]. The same results were obtained when IFN- α -treated macrophages were stimulated with TLR agonists [49]. Furthermore, in a coculture between peripheral blood mononuclear cells and HCV-infected Huh7.5 hepatoma cells, Zhang et al. [23] observed increased IFN- λ secretion after treatment with IFN- α . In conclusion, activation of the type I IFN response clearly primes the cells for IFN- λ production. In IFNAR^{0/0} mice, virus-induced expression of IFN- λ is thus expected to be lower than in wild-type mice. This was indeed observed in IFNAR^{0/0} mice infected with HSV-2, Sendai virus or influenza virus [12, 16].

Less evidence accumulated until now showing that type III IFNs modulate type I IFN expression. Pretreatment of monocyte-derived macrophages or MDDCs with IFN- λ enhanced HSV-1-induced IFN- α and IFN- β mRNA expression [39]. Likewise, IFN- λ treatment of peripheral blood mononuclear cells or pDCs in coculture with HCV-infected Huh7.5 cells enhanced the production of IFN- α [23]. In contrast, two studies failed to show activation of the type I IFN pathway by IFN- λ : after infection with Sendai virus, splenocytes derived from IFN-LR1^{0/0} mice did not produce less type I or type III IFN than splenocytes derived from wild-type mice [12]; also, IFN- λ treatment of human airway epithelial cells infected with influenza A did not increase IFN- β expression [34]. However, these negative results can be interpreted as follows: in the first case, very few cells in the splenocyte population are expected to respond to IFN- λ and therefore to modulate type I IFN expression; in the second case, epithelial cells such as airway epithelial cells are likely poor IFN- β producers, irrespective of priming by IFN.

In conclusion, cross talk exists between the type I and type III IFN systems but it is likely depending on the ability of the cells to produce or respond to specific IFN types.

Potential Therapeutic Use of IFN- λ

Type I IFNs are used to treat diseases such as chronic viral hepatitis or multiple sclerosis. However, these treatments have side effects, which might partly relate to the

Table 3. Association of polymorphism in the region of *IFNL3* with sustained virological response to pegIFN- α and ribavirin therapy

Poly-morphism	Responder allele	Nonresponder allele	Location	Comment	Reference No.
rs12979860	C	T	3 kb upstream of <i>IFNL3</i> , in intron 1 of <i>IFNL4</i>	Strong predictor for HCV clearance	56
rs8099917	T	G	7.5 kb upstream of <i>IFNL3</i>	Strong predictor for HCV clearance	57, 58
rs28416813	C	G	37 bp upstream of <i>IFNL3</i> start codon	In high linkage disequilibrium with rs12979860	56, 68, 69
rs8103142	T	C	<i>IFNL3</i>	In high linkage disequilibrium with rs12979860	56, 68, 69
rs4803217	C	A	3' untranslated region of <i>IFNL3</i>	In high linkage disequilibrium with rs12979860	68, 69
ss469415590	TT	Δ G	<i>IFNL4</i> (exon 1)	Δ G causes expression of IFN- λ 4, which is associated with impaired clearance of HCV Highly correlated with rs12979860[T]	3

Fig. 4. Model of IFN- λ and IFN- α/β responses. By being produced by and acting mostly on epithelial cells, IFN- λ is expected to contribute to an antiviral response that fits anatomical entities such as the gastrointestinal or respiratory mucosae. By acting on most cells, IFN- α/β would act in a rather radial way from the infection focus.

ubiquitous expression of IFNAR. In contrast, the IFN- λ receptor distribution is more restricted [11, 19]. For this reason, fewer side effects may be expected from the use of IFN- λ .

In cultured cells and in the liver, it was shown that repeated stimulation with IFN- α leads to nonresponsiveness of the cells [50, 51]. This IFN- α refractoriness is believed to be one of the reasons for the lack of a response of some HCV-infected patients to the treatment [52]. Interestingly, IFN- λ did not induce such a refractory state in liver cells and might thus be better adapted for repeated treatment [52].

As discussed above, human hepatocytes appear to be responsive to IFN- λ , and clinical trials using recombinant IFN- λ are currently ongoing for the treatment of chronic HCV infection. Phase 1 clinical trials have been published [53, 54]. In these studies, subjects with chronic HCV genotype 1 were administered pegylated IFN (pegIFN)- λ with or without ribavirin. pegIFN- λ was well tolerated and had antiviral activity against HCV. Side effects, such as aminotransferase, lipase or amylase elevation, were reported. Other adverse events, including fatigue, nausea, myalgia or headache, were reported but seemed to be less frequent than after IFN- α therapy. Importantly, no significant hematological toxicity was detected. In a phase 2 study which has only been published as an abstract, the milder side effects of IFN- λ were confirmed [55]. IFN- λ has now entered phase 3 clinical trials.

Genetic Polymorphism at the IFN- λ Locus and HCV Therapy

In 2009, genome-wide association studies have identified single nucleotide polymorphisms (SNPs) in the region the *IFNL3* gene in humans (table 3). These SNPs, rs12979860 and rs8099917, are located ~3 and ~8 kb upstream of *IFNL3*, respectively [56–58], and are associated with response of HCV patients to cotreatment with pegIFN- α and ribavirin, and with spontaneous HCV clearance [56–60]. The rs12979860[T] allele is associated with treatment failure in patients of European ancestry, while the C allele is associated with a twofold greater rate of sustained virological response [56]. Interestingly, Ge et al. [56] identified two additional SNPs in the *IFNL3* gene (SNPs rs28416813 and rs8103142) that are highly associated with rs12979860. rs28416813 consists of a G→C substitution 37 bp upstream of the *IFNL3* start codon, and rs8103142, which occurs in the IFN- λ open reading frame

and results in a nonsynonymous amino acid change at position 70, where arginine is substituted for lysine (K70R). Since then, other SNPs have been identified in the region of the *IFNL3* gene and associated with sustained virological response to pegIFN- α /ribavirin therapy. However, the mechanism by which IFN- λ 3 polymorphisms affect the efficacy of HCV clearance remains to be determined.

Recently, a novel transcript located 3 kb upstream of *IFNL3* was identified as *IFNL4* [3]. The SNP rs12979860 is located within intron 1 of this newly identified gene, and a novel marker, ss469415590, was detected in exon 1. ss469415590 is a dinucleotide variant (TT $\rightarrow$ Δ G) where Δ G results from the deletion of one T nucleotide (rs67272382) and from a T $\rightarrow$ G substitution (rs74597329) [3]. The Δ G variant creates a frameshift that allows the expression of the IFN- λ 4 protein. Paradoxically, the expression of this additional IFN- λ subtype is associated with impaired clearance of HCV, although IFN- λ 4 was shown to exhibit a potent inhibitory effect against HCV replication in vitro [3, 4]. Moreover, IFN- λ 4 expression is often correlated with the expression of the rs12979860[T] allele, which is also a predictor of a low virological response to HCV treatment [3].

Given the association of SNPs in the *IFNL3* locus with spontaneous or drug-mediated viral clearance of HCV, genotyping patients will help to define the therapeutic strategy against this virus.

Concluding Remarks

More and more evidence accumulates showing that the type I and type III IFN systems are largely nonredundant. IFN- λ can induce a longer STAT1 activation and repeated stimulation of cells with IFN- λ does not lead to refractoriness of the cells to subsequent stimulation [52]. Importantly, in vivo, the expression pattern of the IFN- λ receptor clearly diverges from that of the IFN- α / β receptor. The IFN- λ response is predominantly observed in epithelial cells, although some hematopoietic cells were also shown to be responsive. IFN- λ thus likely evolved to protect mucosal surfaces against viral infection. This IFN family is probably of utter importance against viruses with a strong epithelial tropism, as was demonstrated in the case of rotavirus [17]. It is worth noting that recent studies show that epithelial cells also act as IFN- λ -producing cells. The IFN- λ system thus likely evolved to offer a local protection that fits epithelium-rich anatomical structures, such as the gastrointestinal or respiratory tract (fig. 4). Protection of epithelial surfaces may also be instrumental in limiting

virus transmission by feco-oral or aerosol transmission by reducing both virus excretion and virus entry.

The signal transduction pathway leading to IFN- λ production closely resembles that leading to type I IFN production. Yet, some studies suggest that cells producing type I and type III IFNs do not fully overlap and more studies are required to address this issue. Recently, an elegant model of a reporter mouse was developed to identify IFN- β -producing cells. In these mice, one allele of the IFN- β -coding region was replaced by the firefly luciferase-coding region, allowing immunostaining and in vivo imaging of IFN- β -producing cells [61]. The development of such a model to monitor IFN- λ expression in vivo would be particularly valuable to complete the understanding of the peculiarities of the IFN- λ system.

In summary, the characteristics of type III IFNs render them attractive for targeted antiviral therapy. Phase 3 clinical trials are currently ongoing to test the use of IFN- λ 1 in therapy against chronic HCV. Interestingly, a number of polymorphisms have been identified in the region of the *IFNL3* gene and have been shown to influence the outcome of the treatments against HCV. A new IFN- λ subtype, IFN- λ 4, has been described and is implicated in impaired clearance of HCV in patients which express the gene [3]. Understanding the underlying mechanisms of such variations in the responsiveness to treatment will help to improve therapies.

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