Investigations of Interferon-Lambda for the Treatment of Cancer

Andrew Stiff a  William Carson 3rd b

a Biomedical Sciences Graduate Program and b Division of Surgical Oncology, The Ohio State University, Columbus, Ohio, USA

Introduction

While the scientific community has been aware of interferon (IFN) proteins for over 50 years, type III IFNs or IFN-λ were first identified roughly 12 years ago. The IFN-λ lambda family in humans has traditionally consisted of three members: IFN-λ1 (IL-29), IFN-λ2 (IL-28A), and IFN-λ3 (IL-28B) whose genes are all clustered in the q13.13 region of chromosome 19 [1, 2]. These new proteins are related to type I IFN and IL-10 family members [3, 4]. The murine IFN-λ gene cluster is located on chromosome 7A3 and also consists of three genes, but only the IFN-λ2 and IFN-λ3 genes encode functional proteins, whereas the IFN-λ1 gene has lost exon 2 and acquired a stop codon in exon 1 [5]. In addition, a new IFN-λ family member, IFN-λ4, was recently discovered in humans upstream of the gene encoding IFN-λ3. IFN-λ4 was found to be similar to IFN-λ3 and its expression was associated with lower levels of hepatitis C virus RNA prior to treatment, but impaired response to treatment with pegylated IFN-α and ribavirin. Several studies have also identified several polymorphisms within the IFN-λ gene locus of humans, particularly near the IL-28B gene, that have prognostic value in the setting of hepatitis infection [6, 7]. While IFN-λ family members are highly similar to each other in their amino acid sequences, they are only approximately 12 and 15% identical to type I IFNs and IL-10 family members, respectively [1, 2, 8]. Consistent with this, IFN-λ has been found to interact with and sig-
nal through a unique heterodimeric receptor complex consisting of IFN-\(\lambda\) receptor 1 (IFNL1R1) and the IL-10 receptor subunit 2 (IL-10R2) [1–4]. Despite binding to a distinct receptor, IFN-\(\lambda\) has antiviral functions and downstream signaling pathways that overlap significantly with type I IFN [1, 2, 9, 10]. After binding to their respective receptor complexes, both IFN-\(\lambda\) and type I IFNs activate Jak1 and Tyk2 kinases, which phosphorylate STAT1 and STAT2 on tyrosine residues. This leads to STAT1 and STAT2 association with IFN regulatory factor 9 forming the IFN-stimulated gene factor 3 transcriptional complex and the expression of IFN-stimulated genes. IFN-stimulated gene induction by type I IFNs and IFN-\(\lambda\) results in antiviral, antiproliferative, and proapoptotic functions [1, 2, 11, 12]. While these responses are similar, it has been noted that IFN-\(\lambda\) stimulation causes a prolonged activation of STAT1 and STAT2 as well as a delayed but stronger induction of IFN-stimulated genes compared to type I IFN [1, 2, 13]. In the setting of cancer, IFN-\(\lambda\) has been shown to exert direct effects against cancer cells by promoting cell cycle arrest and apoptosis as well as indirect effects through modulation of immune responses [14].

**IFN-\(\lambda\) Tissue Specificity**

Despite the overlapping signaling pathways and antiviral functions performed by IFN-\(\lambda\) and type I IFNs, several important differences have emerged. While the type I IFN receptors IFNAR1 and IFNAR2 are constitutively expressed in the human body, expression of the IFN-\(\lambda\) receptor subunit IFNL1R1 is much more restricted [1, 2]. In an analysis of the expression pattern of IFNL1R1 in human tissues, it was found that IFNL1R1 mRNA levels were highest in the lung, heart, liver, and prostate, while low mRNA levels were detected in the central nervous system, bone marrow, testis, uterus, and skeletal muscle [1, 2, 15]. Within the skin, keratinocytes and melanocytes showed high expression of IFNL1R1 and response to IFN-\(\lambda\), while dermal fibroblasts, endothelial cells and adipocytes did not express IFNL1R1 and were unresponsive to even high doses. High levels of IFNL1R1 and responsiveness to IFN-\(\lambda\) have also been detected in human hepatocytes [1, 2, 15, 16]. Studies in mice have shown a relatively similar pattern of IFN-\(\lambda\) responsiveness in different tissues, with the stomach, intestines, and lungs being the most responsive, while the central nervous system and spleen are relatively insensitive. In particular, it was found that the epithelial cells of most tissues express IFNL1R1 and are responsive to IFN-\(\lambda\) when tested in mice [1, 2, 17, 18].

**IFN-\(\lambda\) and Immune Cell Responses**

The relationship between IFNL1R1 expression and IFN-\(\lambda\) responsiveness is not as straightforward in immune cell subsets. Witte et al. [15] tested the responsiveness of T cells, B cells, monocytes, and NK cells to doses of IFN-\(\lambda\) as high as 1 \(\mu\)g/ml and could detect no response in terms of phosphorylation of STAT1 and STAT3 or increased MHC I expression. The addition of factors known to stimulate these immune cell populations prior to IFN-\(\lambda\) treatment made no difference. While it was found that naïve human B and T cells expressed significant amounts of IFNL1R1, it was determined that they were essentially unresponsive to IFN-\(\lambda\) in terms of STAT1 and STAT3 activation or upregulation of MHC-I expression [1, 2, 15]. It was originally suggested that this discrepancy could be explained by the expression of an IFNL1R1 splice variant (sIFNL1R1) lacking the transmembrane domain. This splice variant was thought to be a soluble form of the receptor that could function as a decoy receptor, thereby negatively regulating IFN-\(\lambda\) signaling in lymphoid tissue [1, 4, 15]. This hypothesis was tested by transiently overexpressing sIFNL1R1 in HepG2 cells. It was found that sIFNL1R1 was secreted into the culture media and could bind IFN-\(\lambda\), thereby inhibiting its ability to induce signaling [1, 15]. In addition, human monocytes and NK cells have also been found to express IFNL1R1, but are unresponsive to IFN-\(\lambda\) [1, 2, 15, 19]. However, conflicting reports have emerged which show that monocytes and T cells are capable of responding to IFN-\(\lambda\) with altered cytokine production. It has been reported that despite a lack of STAT activation or MHC-I expression, monocytes and macrophages do respond to IFN-\(\lambda\) by producing IL-6, IL-8, and IL-10 [20]. In addition, IFN-\(\lambda\) has been found to decrease IL-5 and IL-13 production by naïve and memory human T cells while also increasing production of IFN-\(\gamma\) [20, 21]. This suggests the existence of a STAT-independent signaling network that regulates cytokine production in response to IFN-\(\lambda\) in these cells [1, 2].

**Mechanism of IFNL1R1 Expression**

A recent report investigated the mechanism for tissue specific expression of IFNL1R1 by comparing IFNL1R1 expressing hepatocyte-derived cell lines Huh7 and HepG2 to brain glia-derived cell lines U87 and U373 that do not express IFNL1R1 [22]. These authors found that CpG islands in the promoter region of the IFNL1R1 gene were hypermethylated in the U87 cell line compared to Huh7.
However, treatment of U87 cells with the hypomethylating agent decitabine produced only a minor increase in IFNLR1 mRNA expression, indicating that higher-order chromatin modifications likely also influence IFNLR1 expression. In line with this, they found that Huh7 cells were enriched for markers of open and transcriptionally active chromatin within the IFNLR1 promoter, while U87 cells were enriched for markers of closed and transcriptionally inactive chromatin. Furthermore, treatment of U87 cells with nonspecific histone deacetylase inhibitors (HDACi) and decitabine resulted in significantly increased expression of IFNLR1 in U87 cells as well as cell lines from a number of other tissue types, indicating the response was not tissue specific. Importantly, IFN-λ treatment of U87 cells pretreated with HDACi and decitabine to increase IFNLR1 expression showed increased responsiveness to IFN-λ evidenced by STAT1 phosphorylation, IFN-stimulated gene expression, and antiviral protection. The combination of HDACi and IFN-λ treatment of the U87 glioblastoma and murine B16 melanoma cell line suppressed proliferation and induced apoptosis in these cell lines to a greater extent than either agent alone. These findings suggest that inducing expression of IFNLR1 can be a means to sensitize cells to the anticancer effects of IFN-λ, although other mechanisms of interaction cannot be ruled out given the nonspecific action of HDACi.

### IFN and Cancer Treatment

IFN-α has been extensively studied as an anticancer therapeutic. It has been shown to have antiproliferative, proapoptotic, and immunomodulatory properties in a variety of cancer systems [1, 23–26]. Furthermore, IFN-α has been used extensively in the clinic to treat several malignancies including melanoma, renal cell carcinoma, chronic myeloid leukemia, Kaposi’s sarcoma, and multiple myeloma, among others [2, 23]. In the setting of melanoma, adjuvant therapy IFN-α has produced significant improvements in relapse-free and overall survival in large randomized trials [27, 28]. However, these beneficial effects were only obtained with very high doses (20 IU/m² i.v. 5 doses/week), while studies examining the effects of low-dose IFN-α failed to produce major effects [29, 30]. Such high doses are typically associated with significant side effects such as myelosuppression, neurotoxicity, fever, chills, nausea, and joint pain that sometimes forces the discontinuation of the therapy or lowering of the dose. In many instances these complications reduce the clinical effectiveness of IFN-α therapy, and they are thought to be related to the ubiquitous expression of type I IFN receptors in the human body and their strong effect on immune cells. Given that IFN-λ has significant overlap with type I IFNs in terms of signaling, antiproliferative, and proapoptotic functions, investigations have been initiated into its potential as a new cancer therapy. In addition, given that the profile of IFN-λ responsive cell types is much more restricted compared to IFN-α, it has been hypothesized that IFN-λ could mediate the beneficial aspects of IFN-α with fewer side effects [1, 2].

### Antitumor Effects of IFN-λ

#### Initial Investigation of Anticancer Properties of IFN-λ in Murine Models

Given the clinical use of IFN-α for the treatment of melanoma, the potential to use IFN-λ as a less toxic alternative was pursued. In an initial investigation, Lasfar et al. [5] showed that murine B16 melanoma cells responded to IFN-λ with STAT activation, upregulation of MHC I, and antiviral protection. Subsequently, they transfected B16 cells with a plasmid encoding murine IFN-λ to establish B16 cells that constitutively expressed and secreted murine IFN-λ (B16-IFN-λ). Interestingly, they detected no difference in the proliferation rate between parental B16 cells, control vector-transfected B16 cells, or B16-IFN-λ cells in vitro. They then investigated the potential of constitutive IFN-λ production to affect tumorigenicity in vivo using C57BL/6 mice. In comparison to parental B16 or vector-transfected B16 cells that formed tumors in all mice within 20 days, B16-IFN-λ cells produced tumors significantly later or not at all. In addition, when B16-IFN-λ cells were mixed with equal amounts of B16 parental cells and injected into mice, tumor development was significantly delayed again with a number of mice never forming tumors. To investigate if the decreased tumorigenicity of B16-IFN-λ cells was due to direct effects of IFN-λ on B16 cells or indirect mechanisms such as immune system activation, B16 cells expressing but unresponsive to IFN-λ due to lack of IFNLR1 (B16-IFN-λres) were generated. Similar to B16-IFN-λ cells, B16-IFN-λres cells also demonstrated reduced tumorigenicity, indicating an indirect mechanism likely explained by the effect of IFN-λ. Histological examination showed that B16 parental tumors had enhanced vascularity and a higher mitotic rate than B16-IFN-λ tumors, suggesting a role for IFN-λ in the tumor microenvironment. This group also reported that keratinocytes, which are known to interact
with melanocytes, were highly responsive to IFN-λ. The authors suggested that the response of keratinocytes and other stromal cells in the tumor microenvironment to IFN-λ could play a role in inhibiting tumor growth.

In a subsequent study, Sato et al. [31] generated IFN-λ-expressing murine B16F0 cells (B16F0-IFN-λ). These B16F0-IFN-λ cells showed reduced soft agar colony formation and proliferation in vitro. The authors further demonstrated that B16F0-IFN-λ cells had increased expression of p21 and decreased phosphorylation of Rb, indicating cell cycle arrest. B16F0-IFN-λ cells also showed increased activation of caspase 3 and 7 in vitro, indicating increased apoptosis due to IFN-λ exposure. In addition, B16F0-IFN-λ cells showed reduced pulmonary metastasis following tail-vein injection. This reduction of metastasis was associated with increased cellular infiltration of the lungs, indicating a potential immune response to the malignant cells. In the presence of IFN-λ, depletion of CD4+ and CD8+ T cells did not result in increased B16F0-IFN-λ tumor growth; however, NK cell depletion led to progressive tumor growth, suggesting that NK cells could be the predominant cell type involved in the antitumor action of IFN-λ. A study by Numasaki et al. [32] using a murine fibrosarcoma model provided additional support for an indirect antitumor mechanism of IFN-λ. This study showed that IFN-λ-expressing murine MCA205 cells showed decreased tumor growth and pulmonary metastasis, corroborating previous work. They also showed that sublethal irradiation of the bone marrow was able to abolish this effect, supporting the importance of an immune cell subset in mediating this effect. They went on to show that antibody depletion of NK cells, CD8+ T cells, and neutrophils reduced the effects of IFN-λ, strengthening the argument for an indirect immune-mediated effect.

**Oncolytic Virus Induction of IFN-λ Promotes NK Cell Targeting of Murine Melanoma**

In a study investigating the oncolytic virus vesicular stomatitis virus (VSV), Wongthida et al. [33] hypothesized its mechanism of action against cancer cells in vivo was at least in part mediated by activation of immune cells by the virus that subsequently could target tumor cells. They initially observed that coculture of B16ova cells with bone marrow cells from C57Bl/6 mice alone produce no cytotoxic effects, but B16ova cells were killed when VSV was added to this coculture. This effect occurred with or without the addition of VSV neutralizing serum, which blocks infection and direct oncolysis of B16ova cells by VSV, indicating the virus activated immune cells within the bone marrow cell population that were likely killing the tumor cells. They subsequently found that a B16-derived cell line generated in their lab (B16LIF) was insensitive to VSV-activated BM cells in vitro and in vivo compared to the highly sensitive B16ova in both settings. They found B16ova cells overexpressed IFNLR1 23-fold compared to B16LIF cells, and that B16LIF did not express IFNLR1 even after VSV infection. They went on to show that antibody-mediated blockade of IFN-λ signaling abolished VSV-activated bone marrow cell killing of B16ova cells, suggesting IFN-λ was critical for this function. They were also able to show that depletion of macrophages and Gr1+ cells from bone marrow cell cultures significantly reduced the amount of IFN-λ produced in response to VSV with or without B16ova cells. Depletion of Gr1+ cells and macrophages from bone marrow B16ova cell cocultures also significantly reduced the cytotoxic effect produced by VSV treatment, while depletion of other immune cell subsets had no effect. This study also found that while NK cells were not an important source of IFN-λ in response to VSV, their depletion from B16ova bone marrow cell cocultures reduced the killing of B16ova cells to a similar degree as IFNLR1 blockade. In search of an explanation for this phenomenon Wongthida et al. found treatment of B16ova cells with IFN-λ increased their expression of several NK cell stimulatory ligands, while this did not occur in IFNLR1-deficient B16LIF cells. Furthermore, IFN-λ treatment of B16ova cells was able to induce IFN-γ production by NK cells. These in vitro findings were corroborated in vivo when NK cell depletion significantly reduced the effectiveness of VSV treatment of B16ova tumors. Finally, these researchers used a lentivirus vector to induce expression of IFNLR1 in B16LIF cells, which normally do not express IFNLR1 and are insensitive to VSV-activated bone marrow cell killing or VSV treatment in vivo. Following expression of IFNLR1, B16LIF cells were as sensitive as B16ova cells to VSV treatment in vitro and in vivo, and could effectively activate NK cells following IFN-λ treatment. From this evidence the authors concluded that IFN-λ produced by Gr1+ cells and macrophages in response to VSV treatment led to increased recognition and killing of murine melanoma cells by NK cells. Abushahba et al. [34] corroborated these findings using a murine model in which BNL hepatoma cells were engineered to express IFN-λ. These authors found that while NK cells were unable to directly respond to IFN-λ, tumors expressing IFN-λ showed increased NK cell infiltration, and depletion of NK cells from splenocytes reduced their cytotoxic activity.
**IFN-λ Is Effective Against Human Cancers**

Work done by our group aimed at extending these findings in murine models of melanoma into human systems. As an initial step, mRNA expression of the components of the IFN-λ receptor complex IFNLR1 and IL-10R2 were assessed in eight human melanoma cell lines by RT-PCR. The eight cell lines showed equivalent expression of IL-10R2, but expression of IFNLR1 was variable with 1174 Mel showing undetectable levels. All cell lines tested except 1174 Mel were shown to be responsive to IFN-λ by induction of STAT1 and STAT2 phosphorylation. Interestingly, IFN-λ treatment resulted in STAT5 phosphorylation in 1174 Mel despite it apparently lacking IFNLR1 expression. It was hypothesized that this could occur through IFN-λ’s interaction with other cytokine receptors such as IL-10R1 or IL-20R1. In addition, it was found that IFN-λ treatment did not cause activation of the AKT, ERK, or SAPK/JNK pathways. Microarray gene expression analysis on the 1106 Mel cell line following stimulation with IFN-λ revealed that IFN-stimulated genes such as IFI27, RSAD2, and IFI44L were the most commonly induced genes. IFN-stimulated gene expression was also shown to be dose and time dependent. The potential antitumor mechanisms of IFN-λ against human melanoma cell lines were investigated next. The potential of IFN-λ to prime NK cells and enhance their ability to lyse target melanoma cells was tested. While NK cells were found to express both IL-10R2 and IFNLR1, and showed activation of Jak-STAT signaling in response to IFN-λ, no enhancement of NK cell cytotoxicity was found. The lack of NK cell activity in these experiments was surprising given the evidence supporting the importance of NK cells in murine models. However, these experiments were not conducted in the context of additional immune cells such as macrophages and other lymphocytes, which could play an important role in the murine models. The potential for IFN-λ to exert direct effects against melanoma cells lines was examined next. It was found that IFN-λ treatment had no effect on the proliferation of melanoma cell lines, but that the F01 cell line showed a dose-dependent increase in apoptosis following 48-hour treatment with IFN-λ. It was then shown that the combination of IFN-λ and the proteasome inhibitor bortezomib or IFN-λ in combination with temozolomide was capable of producing synergistic apoptosis in F01 melanoma cells. Finally, the expression of IFN-λ receptor components in benign nevi and primary melanoma lesions was evaluated. All benign nevi examined lacked both IL-10R2 and IFNLR1, while 6 of 8 primary melanoma lesions expressed both components [19].

Other groups have also explored the effects of IFN-λ on a number of different human cancers including bladder carcinoma, Burkitt’s lymphoma, colorectal carcinoma, glioblastoma, non-small cell lung cancer, esophageal carcinoma, and osteosarcoma, among others [1, 2]. In general, these studies have found that cell lines from these various cancers display a range of sensitivity to IFN-λ treatment. This variability generally matches the expression of IFNLR1 by these cell lines, indicating its importance for the effectiveness of IFN-λ treatment of cancer [35–38]. Recently, a report showed that IFN-λ was expressed at higher levels in cervical samples from women infected with low-risk human papilloma virus (HPV) strains compared to high-risk HPV strains. In addition, IFN-λ levels decreased significantly with abnormal cytology, suggesting high-risk HPV strains may be able to counteract the IFN-λ response and that the loss of IFN-λ expression might be related to disease progression [39]. Many of these studies have shown IFN-λ is capable of producing cell cycle arrest in cancer cells through induction of p21 and decreased phosphorylation of Rb as well and the induction of apoptosis evidenced by caspase 3/7 activation or PARP cleavage [31, 36, 37, 40–42]. Several studies have also shown that IFN-λ can increase the effectiveness of several widely used chemotherapeutic agents, suggesting its potential to be used as an adjuvant to existing chemotherapy regimens [19, 36, 37].

**Immune Modulatory Effects of IFN-λ**

Reports investigating the immune modulatory effects of IFN-λ have shown that it is capable of influencing the balance between Th1 and Th2 immune responses [20, 21]. Dai et al. [21] showed that stimulation of human PBMCs with anti-CD2/3/28 beads in the presence of IFN-λ led to a decrease in the production of the Th2 cytokine IL-13 and an increase in IFN-γ production. The authors further showed that IFN-λ led to an immediate decrease in IL-13 production, and it was only at later time points that IFN-γ increased. They concluded from this that IFN-λ acted primarily to inhibit IL-13 production and the increased IFN-γ was a consequence of decreased Th2 cytokine production. They next evaluated the effect of IFN-λ on the production of Th2 cytokines IL-13 and IL-5 by naïve and memory T cells stimulated with anti-CD2/3/28 beads in the presence of IL-4, which promotes the development of a Th2 response. The addition of IL-4 led to a significant increase in IL-13 and IL-5 production by naïve T cells, but did not further increase the produc-
tion of these cytokines by memory T cells, indicating IL-4 had a selective effect on naïve T cells. For both memory and naïve T cells, the addition of IFN-λ led to reduced production of IL-13 and IL-5. They were also able to show that activation of naïve T cells led to an increase in their expression of the IL-4 receptor (IL-4Ra) while this did not occur in memory T cells. Furthermore, the presence of IFN-λ was able to block the induction of the IL-4 receptor on naïve T cells after their activation. Finally, it was found that the addition of IFN-λ to naïve T cells activated with anti-CD2/3/28 beads in the presence of IL-4 led to a significant decrease in expression of the Th2 master transcription factor GATA3, while the same treatments produced only a slight decrease in GATA3 expression by memory T cells. These data led the authors to conclude that IFN-λ was capable of shutting down key pathways involved in Th2 development and therefore allowed the Th1 response to be more robust than would otherwise have occurred in the absence of IFN-λ. It should be noted that the ability of IFN-λ to influence the development of Th1 and Th2 responses stands in contrast to the findings of Witte et al. [15] who found that T cells were unresponsive to IFN-λ treatment. It has been shown that IFN-λ can alter monocyte-derived dendritic cell development such that they diminish Th2 development, suggesting this disagreement could be caused by the effect of IFN-λ on additional immune cell subsets [15, 20].

The potential for IFN-λ’s influence on the balance between Th1 and Th2 immune responses to impact cancer was recently highlighted by an investigation of the IFN-inducible gene USP18 in a mouse model of spontaneous breast cancer by Burkart et al. [43]. USP18 is known to be a negative regulator of type I IFN responses, and USP18 deficiency has been reported to have an inhibitory effect on the development of leukemia. This study used a mouse that was a cross between the polyomavirus middle T (PyVmT) mouse model for breast cancer with a USP18 knockout (KO) mouse to generate PyVmT/USP18 KO mice. In comparison to PyVmT/USP18 WT mice, the PyVmT/USP18 KO mice showed no difference in tumor latency, but did demonstrate a significant improvement in survival and tumor burden. This effect was shown to not be the result of intrinsic differences in proliferation or apoptosis between USP18 KO and wild-type (WT) mammary epithelial cells in this model. Analysis of single-cell suspensions produced from the tumors of PyVmT/USP18 KO and WT mice revealed a significant increase in the number of CD4+ T cells in tumors from USP18 KO mice. There was also an observed increase in CD8+ T cells, NK cells, and macrophages that did not reach statistical significance, while there was no difference in myeloid-derived suppressor cells. Importantly, depletion of CD4+ T cells from FVB WT mice injected with PyVmT/USP18 KO mammary epithelial cells showed enhanced tumor growth, highlighting that reduced tumor growth of USP18 KO tumors was dependent on CD4+ T cells. Interestingly, depletion of CD4+ T cells from FVB WT mice that were injected with both PyVmT/USP18 KO and USP18 WT cells produced a protective effect, indicating USP18 was potentially regulating the pro- and antitumor effects of CD4+ T cells.

To investigate this, Burkart et al. [43] analyzed the expression of T cell-specific chemokines and found that the Cxcr3 ligands Cxcl10 and Cxcl11 were significantly upregulated in PyVmT/USP18 KO tumors. Since expression of Cxcr3 is associated with Th1 cells, the authors investigated if PyVmT/USP18 KO tumors showed a Th1 cytokine profile and found they had increased expression of IFN-γ and decreased expression of the Th2 cytokines IL-4 and IL-13. In addition, when PyVmT/USP18 WT cells were injected into USP18 KO and WT mice, no difference in tumor growth was observed, indicating that it was specifically the lack of USP18 expression in tumor cells that was responsible for this effect. PyVmT/USP18 KO cells stably expressing Cxcl10 shRNA showed increased tumor growth and decreased CD4+ T cell infiltration, highlighting the importance of increased Cxcl10 expression by USP18 KO tumors.

Subsequently, USP18 was identified to be a negative regulator of IFN-λ signaling and Cxcl10 expression in cells with rescued USP18 expression. Finally, injection of PyVmT/USP18 KO mammary epithelial cells with reduced IFNLR1 expression showed increased tumor growth. This work highlights that in the setting of hypersensitivity to IFN-λ signaling due to loss of USP18, the cytokine milieu of the tumor microenvironment is skewed towards a Th1 response that results in reduced tumor growth [43].

**Conclusion and Future Directions**

IFN-λ is a recently discovered cytokine that overlaps significantly with type I IFN in terms of signaling pathways, but differs substantially from type I IFN in terms of tissue responsiveness. This is due to variation in IFNLR1 expression between tissue types, which appears to be epigenetically regulated, potentially allowing for drugs such as HDAC inhibitors to increase cell types that can respond to IFN-λ. IFN-λ has also been shown to exert anti-
tumor effects in both murine and human models. This has been shown to occur through direct effects on target tumor cells as well through indirect immune-mediated responses. IFN-λ can also potentially alter the balance between Th1 and Th2 immune responses, which may be able to be taken advantage of for the treatment of cancer.

Increasing evidence suggests that dysfunction of the immune system plays a critical role in the progression of human malignancies [44]. In fact, some have suggested that evasion of the immune system is in fact a hallmark of cancer [45]. A number of different immune cell subsets including T cells, B cells, NK cells, dendritic cells, macrophages, and myeloid-derived suppressor cells have been identified within the stroma of the tumor microenvironment. These immune cells, as thoroughly reviewed elsewhere [44], are frequently located at distinct locations within the tumor microenvironment, suggesting different functions in controlling or promoting tumor growth. Studies have also shown that an increased lymphocytic infiltrate is associated with an improved prognosis [44].

The ability of IFN-λ to inhibit the development of potentially protumor Th2 immune responses in favor of antitumor Th1 responses suggests it would be interesting to evaluate the efficacy of IFN-λ in combination with emerging immunotherapeutic approaches. In particular, investigation of the ability of IFN-λ to increase expression of potent T cell chemoattractant Ccr3 ligands by human cancers as it does in the murine model discussed above could be productive. This would allow for investigation of the ability of IFN-λ to promote the recruitment of T cells into the tumor microenvironment, and open the door for the investigation of IFN-λ as a complimentary therapy to the new immune checkpoint inhibitors such as ipilimumab (anti-CTLA4) or antibodies directed against PD-1/PD-L1. These immune checkpoint inhibitors function to block inhibitory receptors expressed on T cells whose ligands are often found within the tumor microenvironment or are directly expressed by tumor cells and cause loss of T cell effector function [46, 47]. A common side effect of these checkpoint inhibitors is the development of adverse autoimmune reactions, which suggests their combination with other immune modulatory agents or cytokines could be dangerous. However, as noted above, the limited expression of the IFNLR1 and its lower side-effect profile relative to IFN-α could make IFN-λ a safer alternative in this setting.

References


30 Kleeberg UR, Suciu S, Brocker EB, Ruit DJ, Chartier C, Lienard D, Marsden J, Schadenberg D, Eggermont AM: Final results of the EORTC 18871/DKG 80-1 randomised phase III trial. rIFN-α2b versus rIFN-gamma versus ISD/ASCOR M versus observation after surgery in melanoma patients with either high-risk primary (thickness >3 mm) or regional lymph node metastasis. Eur J Cancer 2004;40:508–515.


