

Type-I IFN Signaling

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The interferon (IFN) family of cytokines is broadly classified into type I, II, and III. The IFN-I consists of 13 subtypes, including IFN α , β , κ , ϵ , ω , and τ , all of which bind to the interferon-alpha receptor complex (IFNAR), composed of the IFNAR1 and IFNAR2 chains.

Keywords: interferon ; inflammation ; autoantibodies

1. Type-I IFN Signaling

Plasmacytoid dendritic cells (DC) are the primary producers of IFN α , and binding of IFN α to IFNAR activates the JAK-STAT signaling pathway that induces interferon-stimulated genes (ISGs) [4]. Together, the JAK-STAT pathway and ISG molecules play crucial roles in the host immune response to bacterial, viral, and fungal diseases and associated inflammation [4]. Pathogen recognition by phagocytes through pathogen-associated molecular patterns (PAMPs) activates multiple and divergent host immune response signaling networks, including the IFN-I response pathways. Subsequent induction of downstream ISGs through the IFN-I pathway facilitates efficient viral killing and controls inflammation. During SARS-CoV-2 infection, the host cytoplasmic molecules, such as retinoic acid-inducible gene-1 (RIG-1) and melanoma differentiation-associated gene-5 (MDA5), recognize intracellular viral RNA and activate the IFN-I pathway, which ultimately enhances the production of effector molecules such as ISGs. Since ISGs regulate a vast range of cellular functions, ranging from transcription to translation of proinflammatory and metabolic molecules in phagocytic cells, any impairment in IFN-I signaling can be harmful to the host. Thus, optimal and timely activation of IFN-I signaling is crucial for a well-coordinated antiviral effect of phagocytes with minimal damage to the host [1][2][3][4].

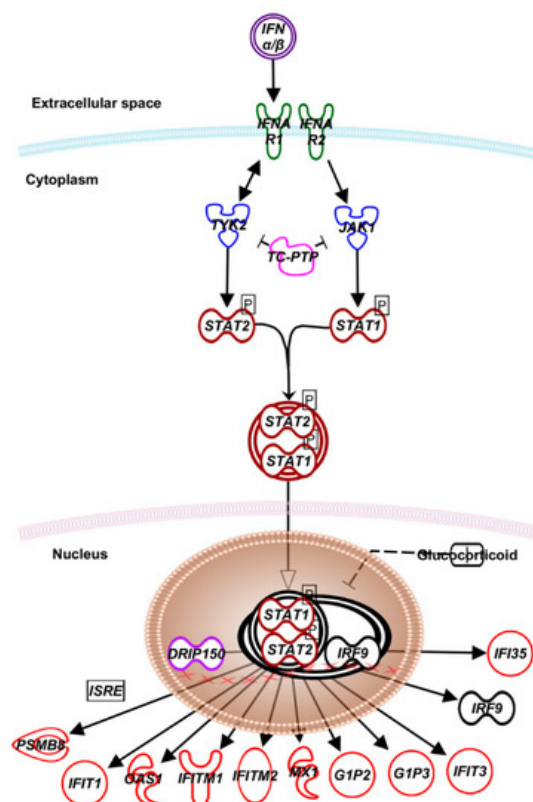


Figure 1. Canonical type I interferon signaling. IFN-I family of cytokines, encompassing IFN- α , - β , and - κ play critical roles in initiating, maintaining, and regulating host immune responses, particularly against viral infections [5][6]. The IFN signaling starts with type I and II IFN receptors (IFNAR), which are transmembrane proteins that function as heterodimers. While type I IFNAR comprises AR1 and AR2 subunits, type II IFNAR is made of GR1 and GR2. The IFNAR1 and IFNGR1

are involved in binding type I and type II IFNs, respectively, while IFNAR2 and IFNGR2 subunits are involved in signal transduction. Engagement of IFNAR2 and IFNGR2 activates tyrosine kinases JAK1, JAK2, and TYK2 that phosphorylate the STAT1 and 2 proteins. After dimerization, the STAT proteins translocate to the nucleus. In the case to type I IFNs, the STAT1/2 heterodimers associate with IFN gene regulatory factor-9 (IRF-9), forming ISGF3, a trimeric transcription factor. The vitamin D receptor-interacting protein150 (DRIP150) interacts with STAT2 and stimulates IFN-dependent transcription. ISGF3 and DRIP150 complex bind to the IFN-stimulated regulatory elements (ISREs) to induce the expression of several IFN- α/β -regulated genes [5][6]. Abbreviations: TYK2—tyrosine kinase 2; JAK1—Janus kinase 1; TC-PTP—T-cell protein tyrosine phosphatase; STAT—signal transducer and activator of transcription; IFI—interferon-induced protein; DRIP150—vitamin D receptor-interacting protein 150; IRF—interferon regulatory factor; PSMB—proteasome 20S subunit beta; IFIT—interferon-induced protein with tetratricopeptide repeats; OAS—O-acetyl serine; MX1—MX dynamin-like GTPase 1; G1P2—ISG-15 ubiquitin-like modifier; ISRE—interferon-stimulated response element.

Apart from ISGs, IFN-I augments host protection against viral infections by recruiting and activating innate (DC, natural killer (NK) cells, neutrophils, monocytes, and macrophages) and adaptive (T and B) immune cells [5][6].

2. Type-I IFN Response and IFN-I Immunotherapy for COVID-19

The IFN-I pathway also plays a central role in the pathogenesis of COVID-19 [4]. Perturbations in IFN-I and ISG levels are among the primary indicators of disease severity in COVID-19 cases. A significant reduction in serum IFN-I levels has been reported in critically ill COVID-19 patients with a high viral load in their respiratory tract than those with mild or moderate disease symptoms. Consistently, the number of plasmacytoid DC, the primary producers of IFN α , was reduced in patients with severe COVID-19, contributing to reduced IFN-I levels. However, data from SARS-CoV-2 infection studies conducted on lung epithelial cells expressing angiotensin-converting enzyme-2 (ACE2) receptor, as well as in human organoids, human bronchial cells, and in post-mortem lung samples of COVID-19 cases showed a moderate-to-high level of ISGs (e.g., MX1, MX2, OAS1, IFI6, IFI27, IFIT1, IFIT3, IFITM1, IFITM2, and IFITM3) despite a low-to-undetectable IFN-I level. This observation suggests that either the infected host cells are refractory to the COVID-19 defense mechanism or the uninfected cells also contribute to elevated ISG level (bystander effect).

Induction of ISGs in infected cells mounts a robust antiviral response against SARS-CoV-2, including blocking viral replication. Besides, upregulation of ISGs, such as IFIT and IFITM family genes, including OASL, OAS2, IFI6, MX1, and MX2, as well as proinflammatory cytokines, including IL-1 β IL-10, TNF- α , IL-6, and IL-1Ra was noted in the bronchoalveolar lavage fluid of COVID-19 patients [7][8][9]. The expression level of several proinflammatory genes involved in IFN-I signaling, including MX1, STAT1, STAT2, IFITM1, IFIT2, IFI35, IFIH1, IRF7, and IFI16 was enriched in COVID-19 patients [10]. Furthermore, these molecules' expression patterns could differentiate mild/moderate COVID-19 cases from healthy individuals [10].

References

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2. Khalaf, K.; Papp, N.; Chou, J.T.; Hana, D.; Mackiewicz, A.; Kaczmarek, M. SARS-CoV-2: Pathogenesis, and number of patient samples used in gene expression studies: it is premature to conclusively define biomarkers of disease severity in COVID-19, based purely on the expression pattern of a specific set of genes. Thus, extensive profiling of enriched inflammatory gene expression from different clinical samples in many patients with various disease severity is required to define and correlate biomarkers with corresponding prognoses of COVID-19 cases. A recent study on the transcriptome of human bronchial cells, and the pulmonary organoids of non-human primates reported that ACE2, the primary receptor for SARS-CoV-2 entry into host cells, is also a member of ISG. This study also indicated a dual role for IFN-specific upregulation of ACE2 in tissue tolerance and infection upon SARS-CoV-2 entry into host cells [11]. Although 5. Lee, A.J.; Ashkar, A.A. The Dual Nature of Type I and Type II Interferons. *Front. Immunol.* 2018, 9, 2061.
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Comorbidities and genetic susceptibility of the host can also alter the IFN-I response to viral infections and cause detrimental effects to the host [10]. During the initial stages of COVID-19, when the viral load, disease severity, and antibody responses are low, IFN-I plays a beneficial host effect by inhibiting viral propagation in the host. Consistently, elevated levels of IFN-I were observed at the infection site and in the blood of COVID-19 cases with mild symptoms and low viral burden. Thus, elevated IFN-I appears to be a host response against SARS-CoV-2 replication, dampening infection and alleviating disease symptoms. However, in COVID-19 patients with severe disease symptoms and a higher viral load, including the elderly population and those with comorbid conditions, the expression of IFN-I was significantly suppressed, and was associated with exacerbated tissue inflammation and disease pathology [12].

Notably, the causal link between the impaired type I IFN response and the severity of COVID-19 remains unknown. Similarly, the net effect of IFN-I-mediated antiviral and anti-inflammatory activities during COVID-19-mediated tissue damage is yet to be determined.

- [illegible]

Trial ID	IFN-I Molecule	Adjunctive Drug #	Route of IFN-I Administration	Study Sponsor	Reference/Status *
2020-000936-23 (DisCoVeRy Trial)	IFN β 1 α	Standard care	Subcutaneous	INSERM, France	Recruiting
NCT04315948 (DisCoVeRy Trial)	IFN β 1 α	Lopinavir/Ritonavir or Standard care	Subcutaneous	Institut National de la Sante Et de la Recherche Medicale, France	Recruiting
NCT04276688	IFN β 1 β	Lopinavir/Ritonavir, Ribavirin	Subcutaneous	The University of Hong Kong, Hong Kong.	Recruitment completed
NCT04254874	IFN α 2 β	Abidol hydrochloride	Atomization (nasal spray)	Tongji Hospital, China	Recruiting
NCT04293887	IFN α 1 β	Lopinavir/Ritonavir, Remdesivir	Nebulization inhalation	Tongji Hospital, China	Not recruiting
NCT04552379	IFN β 1 α	Standard care	Subcutaneous	Pontificia Universidad Catolica de Chile	Recruiting
NCT04273763	IFN α 2 β	Bromhexine Hydrochloride, Abidol hydrochloride and Standard care	Atomization (nasal spray)	WanBangDe Pharmaceuticals, China	Not recruiting
NCT04291729	IFN α 2	Danoprevir/Ritonavir	Spray inhalation	The Ninth Hospital of Nanchang, China	Completed
NCT04251871	IFN α	Lopinavir/Ritonavir or Traditional Chinese Medicines	Aerosol inhalation	Beijing 302 Hospital, China	Recruiting
NCT04275388	IFN α	Lopinavir/Ritonavir and Abidor Hydrochloride	Nebulization inhalation	Jiangxi Qingfeng Pharmaceutical Co. Ltd.	Not recruiting
NCT04469491	IFN β 1 β	Standard care	Nebulization inhalation	Centre Hospitalier Universitaire, Amiens	Suspended

\$ Data obtained from the WHO COVID-19 Vaccine website [1]. * status date was current as of 3 January 2021. # some studies involve “Standard care”, which includes treatment with/without antibiotics, corticosteroids or hydroxychloroquine and oxygen support, in addition to IFN-I therapy. ChiCTR—Chinese Clinical Trial Registry; IRCT—Iranian Registry of Clinical Trials; RECOVERY—Randomized Evaluation of COVID-19 Therapy; DisCoVeRy—Trial of Treatments for COVID-19 in Hospitalized Adults; NCT—National Clinical Trial.

3. Implications of IFN-I Immunotherapy on the Host Response to Other Infectious Diseases

Below is a summary of health conditions in which IFN-I immunotherapy can potentially impact the host response to bacterial and viral infections and the clinical outcome of treatment.

3.1. Viral Diseases

IFN therapy has been credited as prophylaxis to prevent virus-induced exacerbations in airway diseases, such as chronic obstructive pulmonary disease (COPD). A recent report using the primary cell culture model of respiratory syncytial virus (RSV) and influenza virus infection shows that intermittent treatment with IFN- β was beneficial to the host cells and did not trigger proinflammatory cytokines, such as IL-1 β , TNF, and IL-6 [13]. This observation suggests that IFN-I therapy helps to prevent COVID-19-associated pulmonary exacerbations in patients with pre-existing COPD or co-infected with other viral pathogens such as RSV.

The role of IFN-I in human immunodeficiency virus (HIV) infection and its therapeutic role is inconclusive. Upon the entry of HIV-1, the pathogen recognition receptors (PRR) on phagocytes, including Toll-like receptor-7 (TLR7), cGAS, and IFI16, recognize the viral antigens and secrete IFN-I. The binding of IFN-I to IFNAR and phosphorylation of the downstream JAK/STAT pathway leads to the upregulation of ISGs, which are proven antiviral defense molecules (Figure 1). Since the induction of ISGs can deplete CD4+ and CD8+ cells, a low level of IFN-I during the early stages of HIV-1

infection was presumed beneficial. Thus, a delayed and/or sub-optimal IFN-I production can lead to an immunosuppressive niche and persistent, chronic disease. However, during latent HIV-1 infection, sustained IFN-I signaling is associated with immune activation and viral persistence. In summary, IFN-I signaling has a paradoxical yet dynamic role in the pathogenesis of acute vs. chronic/persistent HIV-1 infection ^{[14][15]}. Hence, caution should be taken when administering IFN-I treatment to COVID-19 patients who are also HIV-positive. Clinical parameters, such as blood CD4+ T cell counts and co-infection with other pathogens, should be monitored in these patients during IFN therapy.

3.2. Nosocomial Infection

Reports have shown that COVID-19 cases acquire secondary nosocomial infections, including multidrug-resistant *Staphylococcus aureus* and *Streptococcus pneumoniae*, possibly due to lower IFN-I levels in these patients. However, excessive IFN-I levels have been associated with increased susceptibility to disease progression with worsened pneumonia. Moreover, elevated IFN-I molecules inhibit the proinflammatory TH17 signaling and impair the recruitment of macrophages and neutrophils, as well as neutrophil chemokine secretion, all of which facilitate the progression of nosocomial infection ^[16]. However, the correlation between IFN-I levels and disease severity in COVID-19 or secondary nosocomial infections is not fully understood. Since critically ill COVID-19 patients in hospitals are more prone to nosocomial infections, the dose and route of administration of IFN-I and therapy duration should be planned cautiously.

3.3. Tuberculosis

Upregulation of IFN-I pathway molecules has been reported during infection by *Mycobacterium tuberculosis* (Mtb), the causative agent of tuberculosis (TB), a contagious bacterial disease marked by severe, chronic lung inflammation ^[17]. TB, primarily a pulmonary disease, caused 1.5 million deaths worldwide and an estimated 10 million new cases in 2018 (https://www.who.int/tb/publications/factsheet_global.pdf?ua=1). Elevated IFN-I levels correlate with the disease severity caused by hypervirulent Mtb strains and increased host susceptibility to active TB. Several possible mechanisms, including suppression of proinflammatory innate and Th1 responses, early cell death of alveolar macrophages, recruitment of permissive myeloid cells, and the reduced macrophage antibacterial responses, are associated with the detrimental effect of elevated IFN-I in Mtb-infected host ^[17]. However, when co-administered with anti-mycobacterial therapy or Bacille Calmette-Guerin (BCG) vaccine, IFN- α conferred host-protective effect against Mtb infection. Studies using human PBMCs and animal models showed that priming with IFN- α increases the BCG vaccine's efficacy and promotes T-cell priming and IFN- γ -induced production of "host-protective" proinflammatory cytokines, including IL-12, TNF- α , IL-17, and IL-1 β ^[18]. Notably, the role of IFN-I in regulating chronic, asymptomatic latent Mtb infection (LTBI) and their potential for reactivating LTBI to active TB disease remains unclear. Thus, IFN-I treatment of COVID-19 cases with underlying LTBI might lead to TB reactivation. Moreover, patients with symptomatic, active TB have elevated IFN-I levels ^[17]. Therefore, in the context of Mtb infection, the effect of IFN-I response can be beneficial or detrimental to the host, dependent on whether the host is immunized with BCG or having LTBI/active TB, respectively. In this regard, measuring the IFN-I level in samples, and TB or LTBI status of the patient, determined by skin test or quantiferon test or chest X-ray, would be valuable for initiating IFN therapy for COVID-19 cases. This approach would alleviate the risk for reactivation of LTBI or exacerbation of TB among COVID-19 patients.