

ISG15 regulates IFN- γ immunity in human mycobacterial disease

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Interferon-gamma (IFN- γ) is crucial for immunity against different pathogens due to its broad effects on the multiple arms of the immune system. The regulation of IFN- γ immunity is of extensive interest to research as well as practical activity for drug discovery. New evidence supports previous findings that ubiquitin-like protein ISG15 acts as an extracellular cytokine and promotes IFN- γ production, providing intriguing insights of the importance of ISG15 into the control of human mycobacterial disease.

ISG15 is an interferon-stimulated gene (ISG) [1]. It is strongly upregulated by interferons (IFNs), especially type I IFN (IFN- α/β), pathogen infections, and cellular stresses that activate IFN production [2]. ISG15 is the first identified ubiquitin-like modifier, which can covalently conjugate to other cellular proteins to form ISGylated proteins. Free ISG15 is detected as both extracellular and intracellular protein. The mature form of ISG15 contains two ubiquitin-like domains with a calculated molecular weight of 17 145 Daltons. It lacks an N-terminal methionine, and has a C-terminus ending with the amino acid sequence 'Leu Arg Leu Arg Gly Gly' (LRLRGG), which is the same six amino acid sequence as mature ubiquitin. As with the ubiquitin system, there are a series of distinct enzymes involved in the process of protein ISGylation, including ISG15 activating enzyme (E1), UBE1L — conjugating enzyme (E2), UBCH8 — protein ligases (E3), and ISG15-specific protease

USP18. However, unlike ubiquitin and many ubiquitin-like proteins, such as SUMO1 and NEDD8, which can be found in all species of eukaryotic cells, ISG15 is only identified in vertebrates, including human, monkey and mouse, *etc.* Moreover, compared with ubiquitin, which has virtually 100% cross-species conservation, ISG15 protein has relatively low cross-species conservation, ranging from a high of 98% (chimpanzee to human) to a low of 42% (opossum to human) in mammals. Together, the relatively low cross-species conservation and the absence in many eukaryotic species indicate that *ISG15* is not an essential housekeeping gene but may be an evolutionary adaptation to particular conditions [2].

Significant efforts have been made to identify the substrates of ISG15 conjugation and their roles in pathogen infection and tumorigenesis [3, 4]. For example, the molecular basis of antiviral effect of ISG15 was described in a recent report of the influenza A viral protein NS1 (NS1A). Multiple lysines of the NS1A protein could be modified by ISG15, with K41 in the N-terminal domain of this protein acting as the main modification site. ISG15 modification on NS1A resulted in a disruption of normal function of NS1A, thus inhibiting the virus infection [3]. Meanwhile, a recent report revealed an important role of ISG15 in the enhancement of chemosensitivity during cancer therapy [4]. Treatment of cells with several chemotherapeutic drugs, including doxorubicin and camptothecin, increases ISG15 expression and results in ISGylation of

Δ Np63 α , an alternative splice variant of p53 family protein p63. The ISGylated nuclear Δ Np63 α is subjected to the cleavage by caspase-2 and subsequently releases its inhibitory domain to the cytoplasm, resulting in the inactivation of Δ Np63 α . Since Δ Np63 α inhibits the activity of p53 family members in a dominant-negative manner to increase cell proliferation and tumorigenesis, ISGylation of Δ Np63 α ablated the ability of Δ Np63 α to promote anchorage-independent cell growth and tumor formation *in vivo*.

Besides protein ISGylation, significant amount of free ISG15 was also detected in both intracellular and extracellular compartments. Several studies suggested a potential role of free ISG15 as a cytokine or chemokine [5-7]. An early study in 1996 demonstrated that free ISG15 is secreted from multiple cells, including primary cultures of peripheral blood CD3⁺ cells and viable cell lines of monocyte, T lymphocyte, B lymphocyte, and epithelial origins, and is detectable in serum from healthy human volunteers treated with IFN- β [6]. The potential modulation of immune cell function by ISG15 was reported later in the same year. ISG15 stimulates the production of IFN- γ from CD3⁺ T cells to enhance the proliferation and cytotoxicity of natural killer (NK) cells [5]. In the year of 2003, ISG15 was shown as a neutrophil chemotactic factor in *Plasmodium yoelii*-infected red blood cells [7]. However, a primary challenge to the cytokine properties of ISG15 is that cytokine- or chemokine-related defects have not been identified

rate of ~40% at 45 weeks post infection compared to almost no death in the control mouse group within the same period. These data strongly indicate the essential role of ISG15 in anti-mycobacterial disease.

In summary, the discovery by Bogunovic and colleagues has revealed the critical role of ISG15 in human health. It provides new evidence for ISG15-regulated IFN- γ immunity, which is summarized in Figure 1. The report also clearly shows ISG15 cytokine-related defects in *ISG15* knockout mice, the lack of which has been challenging the idea that ISG15 acts as a cytokine for over a decade. These findings may also answer the question why ISG15 is only induced by IFNs or pathogen infection but not constitutively expressed to counter potential pathogen infection. Possibly, constitutive ISG15 expression and active IFN- γ immunity is deleterious in contexts other than mycobacterial infection. For example, in certain cases of autoimmune disease, a blockade of IFN- γ immunity actually limits the diseases [12, 13]. The cytokine properties

of ISG15 are reminiscent of potential cytokine properties of ubiquitin and other ubiquitin-like proteins. Several studies have suggested that extracellular ubiquitins serve as immune modulator during infectious and noninfectious inflammation [14, 15]. However, not many reports are available regarding other ubiquitin-like proteins. Meanwhile, future investigation focusing on how ISG15 is secreted from certain cells, which cell surface receptors and cellular pathways are responsible for ISG15-induced IFN- γ production, and how the components of these pathways are targeted to regulate IFN- γ immunity, will turn out to be fruitful and point to more applicable strategies for ISG15-targeted drug discovery.

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