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Negative regulation of protein phosphatase 2Cβ by ISG15 conjugation

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Abstract

ISG15, an interferon-upregulated ubiquitin-like protein, is covalently conjugated to various cellular proteins (ISGylation). In this study, we found that protein phosphatase 2Cβ (PP2Cβ), which functions in the NF-κB pathway via dephosphorylation of TAK1, was ISGylated, and analysis by NF-κB luciferase reporter assay revealed that PP2Cβ activity was suppressed by co-expression of ISG15, UBE1L, and UbcH8. We determined the ISGylation sites of PP2Cβ and constructed its ISGylation-resistant mutant. In contrast to the wild type, this mutant suppressed the NF-κB pathway even in the presence of ISG15, UBE1L, and UbcH8. Thus, we propose that ISGylation negatively regulates PP2Cβ activity.

Keywords: ISG15; Interferon; Ubiquitin; Protein phosphatase; NF-κB

Abbreviations: ISG15, interferon-stimulated gene 15 kDa; E1, ubiquitin-activating enzyme; E2, ubiquitin-conjugating enzyme; PP2Cβ, protein phosphatase 2Cβ; TAK1, TGF-β-activated kinase; TAB1, TAK1-binding protein; NF-κB, nuclear factor κB; IκB, inhibitor of NF-κB; IKK, IκB kinase; IFN, interferon
1. Introduction

Interferon (IFN) is a pleiotropic cytokine that has an essential role in cellular antiviral response through inducing interferon-stimulated genes (ISGs) [1]. ISG15, one of the ISGs, is a ubiquitin-like protein that is covalently conjugated to various cellular proteins (ISGylation) upon interferon stimuli [2]. This protein functions as an antiviral protein against Sindbis virus and HIV-1 [3,4] and as a suppressor of the ubiquitin-proteasome system [5]. Although various target proteins for ISGylation have been identified by a proteomic approach [6-8] and a cascade of the protein ISGylation system has become clear [9-14], the biological consequences of ISGylation of target proteins have been studied in only a few cases [15-17].

Protein phosphatase 2Cβ (PP2Cβ) is an enzyme that belongs to the PP2C type protein phosphatase family and functions as a monomer [18]. PP2Cβ dephosphorylates and suppresses TAK1 and IKK, both of which have essential roles in the NF-κB pathway, an important pathway functioning in innate immunity, adaptive immunity, and cancer [19-22].

We previously reported that ISGylation of UbcH6 and Ubc13 suppresses their ubiquitin E2 enzyme activities [16,17]. In a search for ISGylated proteins, we identified PP2Cβ as a target protein for ISGylation. In addition, we found that ISGylation of PP2Cβ suppresses the activity of PP2Cβ against TAK1-induced NF-κB activation.

2. Materials and methods
2-1. Cell culture and transfection

HeLa cells were cultured in Dulbecco’s modified Eagle’s medium (Sigma) supplemented with 10% heat-inactivated calf serum (Hyclone). Transfection was performed according to the standard calcium precipitation protocol.

2-2. Plasmid construction

The mammalian expression plasmids of ISG15, UBE1L, and UbcH8 were generated as described previously [17]. The open-reading frames of human PP2Cβ, TAK1, and TAB1 were amplified by PCR. Deletion and point mutants of PP2Cβ were generated by PCR. All constructs were verified by DNA sequencing. To generate mammalian expression plasmids of C-terminally and N-terminally Flag-tagged PP2Cβ, the PCR fragment was subcloned into pCI-neo-C3Flag and pCI-neo-3Flag vectors, respectively, which had been generated by inserting three repeats of Flag tag sequence into the pCI-neo mammalian expression vector (Promega). To generate the mammalian expression plasmids of TAK1 and TAB1, the PCR fragments were subcloned into the pCI-neo-2S and pCI-neo-5HA vectors that had been generated by inserting oligonucleotides encoding two repeats of S peptide sequence and five repeats of HA tag sequence, respectively, into the pCI-neo mammalian expression vector.
2-3. Immunoprecipitation and Western blotting

HeLa cells that had been transiently transfected with indicated plasmids and cultured for 24-30 h were lysed with RIPA buffer containing 50 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.1% SDS, 0.5% sodium deoxycholate, 1% Nonidet P-40, 1 mM phenylmethylsulfonyl fluoride and 5 mM N-ethylmaleimide, and the supernatant of the cell lysate was subjected to immunoprecipitation and then to Western blotting with various antibodies as described previously [16,17] except for anti-PP2Cβ [19] and anti-IκBα (BD Biosciences) antibodies. The effect of IFN treatment was analyzed as described previously [17] except for HeLa cells and lysis buffer containing 20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1% Triton X-100, 5 mM EDTA, 1 mM phenylmethylsulfonyl fluoride, and 5 mM N-ethylmaleimide.

2-4. Reporter gene assay

HeLa cells were transfected with the pNF-κB-Luc reporter plasmid (BD Biosciences) and indicated plasmids, together with pRL-TK (Promega) for normalizing transfection efficiency. Twenty-four h after transfection, cells were lysed and luciferase activity was measured by using a Dual-Luciferase Reporter Assay System (Promega) and an AB-2000 luminescence-PSN (Atto, Tokyo, Japan). The same experiments were repeated three times.

3. Results and discussion
To identify proteins modified with ISG15, we carried out a proteomic analysis of ISGylated proteins. We expressed Flag-tagged ISG15, S-tagged UBE1L (E1), and S-tagged UbcH8 (E2) in HeLa cells, and ISGylated proteins were isolated by immunoprecipitation with anti-Flag antibody and subjected to peptide mass fingerprinting. We identified several novel ISGylated proteins (data not shown; data to be presented elsewhere). Among them, we focused on PP2Cβ because PP2Cβ dephosphorylates and suppresses TAK1 and IKK, kinases both functioning in the NF-κB pathway [19,20].

To confirm ISGylation of PP2Cβ, we expressed Flag-tagged PP2Cβ together with T7-tagged ISG15, S-tagged UBE1L and S-tagged UbcH8 in HeLa cells, and the extract of transfected cells was subjected to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-T7 tag and anti-Flag tag antibodies (Fig. 1). Two bands (the molecular masses of 88 kDa and 110 kDa) with slower mobilities than that of intact PP2Cβ (67 kDa) were detected by anti-Flag tag antibody in UBE1L- and UbcH8-dependent manners and these bands were also detected by anti-T7 tag antibody: The above difference in molecular masses between modified and intact ones is reasonably thought to be due to ISGylation. These results strongly suggest that PP2Cβ is covalently modified with ISG15, the conjugation being mediated by UBE1L and UbcH8. It should be noted that ISGylation of
various proteins was detected by over-expression of UBE1L in the presence of ISG15 and was further enhanced by co-expression of UBE1L and UbcH8 (Fig. 1, left panel): These results are consistent with the previous report [11].

Next, to confirm that PP2Cβ is ISGylated in response to IFN signal, HeLa cells that had expressed Flag-tagged PP2Cβ were treated with IFNβ. The cell extract was subjected to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-Flag tag and anti-ISG15 antibodies (Fig. 2A). In another experiment (Fig. 2B), the extract of HeLa cells that had been treated with IFNβ was subjected to immunoprecipitation with anti-PP2Cβ antibody and then to Western blotting with anti-PP2Cβ and anti-ISG15 antibodies. In both experiments, two bands due to ISGylated PP2Cβ were detected only in the case of IFN treatment, indicating that PP2Cβ is ISGylated in response to IFN signal under physiological conditions. ISGylation of various proteins was also detected only in the case of IFN treatment (Fig. 2A and B, right panels). It should be noted that the molecular mass (67 kDa) of exogenously expressed Flag-tagged PP2Cβ isoform 1, the longest isoform of human PP2Cβ [23] and the isoform used in this study, is different from that (42 kDa) of endogenous PP2Cβ isoform 2, the most abundant isoform in HeLa cells [24].

3-2. PP2Cβ activity is suppressed by addition of a protein ISGylation system

It has been reported that PP2Cβ dephosphorylates and suppresses TAK1 [19], which
functions in the NF-κB pathway together with TAB1 [25]. We therefore carried out an experiment to determine whether a protein ISGylation system modulates PP2Cβ activity against TAK1/TAB1-induced NF-κB activation. HeLa cells were co-transfected with plasmids of HA-tagged TAK1 and TAB1, Flag-tagged PP2Cβ, and ISGylation system (T7-tagged ISG15, S-tagged UBE1L, and S-tagged UbcH8), together with NF-κB luciferase reporter plasmid, and luciferase activity was measured (Fig. 3). PP2Cβ suppressed TAK1/TAB1-induced NF-κB activation in the absence of the ISGylation system but not in the presence of the ISGylation system. It should be noted that TAB1 alone has little effect [25]. These results suggest that ISGylation of PP2Cβ suppresses the activity of PP2Cβ against TAK1/TAB1-induced NF-κB activation.

3-3. PP2Cβ is modified with ISG15 at least through Lys12 and Lys142

To confirm the above assumption, it is necessary to construct an ISGylation-resistant mutant of PP2Cβ. Since PP2Cβ is expected to be ISGylated through at least two Lys residues (see Fig. 1), we first constructed various PP2Cβ deletion mutants and expressed them together with the above ISGylation system in HeLa cells. The extract of transfected cells was subjected to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-Flag tag and anti-T7 tag antibodies (Fig. 4). PP2Cβ Δ296-479 was modified with ISG15 to an extent similar to that in the case of the wild type, while PP2Cβ Δ1-295 was not
modified with ISG15 (Fig. 4A) and PP2Cβ Δ1-92 and Δ93-193 seemed to be less modified (Fig. 4B). We hypothesized that one of the Lys residues modified with ISG15 is located in the sequence 1-92, while another is in 93-193. We next constructed various PP2Cβ mutants in which the Lys residues had been changed to Arg, measured their levels of ISGylation, and found that PP2Cβ K12R was less modified with ISG15 than was the wild type (data not shown). We finally constructed various PP2Cβ KR mutants in which the two Lys residues had been changed to Arg and analyzed their levels of ISGylation (Fig. 5). PP2Cβ K12,142R was barely modified with ISG15 in either the case of N-terminally (Fig. 5A) or C-terminally Flag-tagged PP2Cβ (Fig. 5B). Thus, we concluded that PP2Cβ is modified with ISG15 at least through Lys12 and Lys142.

3-4. PP2Cβ activity is suppressed by ISG15 conjugation

Using the PP2Cβ K12,142R mutant as an ISGylation-resistant mutant, we performed an NF-κB luciferase reporter assay as described above (see Fig. 3). As shown in Fig. 6A, in the absence of the ISGylation system, the wild type (Wt) and the K12,142R (KR) mutant suppressed TAK1/TAB1-induced NF-κB activation, while in the presence of the ISGylation system, the wild type had little suppressive effect as shown above (see Fig. 3), but the K12,142R mutant had a suppressive effect. It should be noted that the ISGylation system alone has little effect (data not shown). These results suggest that ISGylation of PP2Cβ
suppresses its activity.

Since degradation of IκBα by the ubiquitin-proteasome system is essential for activation of NF-κB [26-28], we next carried out an experiment to clarify the effect of ISGylation of PP2Cβ on IκBα degradation. We expressed Flag-tagged wild-type PP2Cβ or its K12,142R mutant, S-tagged TAK1 and TAB1, and the ISGylation system in HeLa cells, and the cells were then treated with cycloheximide (CHX), a protein synthesis inhibitor, for 30 min. The cell extract was subjected to Western blotting with anti-IκBα, anti-Flag tag, anti-S peptide, and anti-T7 tag antibodies (Fig. 6B). In our assay conditions, in the absence of cycloheximide, the amount of IκBα protein remained unchanged in all combinations of expression plasmids. However, when the cells had been treated with cycloheximide, the level of IκBα in the case of wild-type PP2Cβ was less than that in the case of the K12,142R mutant in the presence of the ISGylation system. It should be noted that the ISGylation system alone has little effect (data not shown). These results imply that ISGylated PP2Cβ has a less suppressive effect on TAK1/TAB1-induced degradation of IκBα than the ISGylation-resistant mutant does, supporting the above assumption that ISGylation of PP2Cβ suppresses its activity.

In conclusion, we found that PP2Cβ is covalently modified with ISG15 in an IFN-dependent manner and that ISGylation of PP2Cβ suppresses the activity of PP2Cβ against TAK1/TAB1-induced IκBα degradation and NF-κB activation. It can be inferred that the activity of ISGylated PP2Cβ is less than that of the original one. The
construction of an in vitro assay system will clarify this point and is our future project.

Considering that TAK1 is indispensable for cellular responses to various stimuli, including interleukin-1, tumor necrosis factor-α, and a toll-like receptor ligand [25,29], it can be inferred that ISGylation of PP2Cβ has an effect on various cellular responses.

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References


interferes with the ubiquitin/26S proteasome pathway. Cancer Res. 66, 921-928.


ISG15 conjugation targets both IFN-induced and constitutively expressed proteins 
functioning in diverse cellular pathways. Proc. Natl. Acad. Sci. USA 102, 
10200-10205.

[8] Giannakopoulos, N.V., Luo, J.K., Papov, V., Zou, W., Lenschow, D.J., Jacobs, B.S., 
Commun. 336, 496-506.


Huibregtse, J.M. and Krug, R.M. (2004) The UbcH8 ubiquitin E2 enzyme is also the


between the ubiquitin conjugation system and the ISG15 conjugation system: ISG15 conjugation to the UbcH6 ubiquitin E2 enzyme. J. Biochem. (Tokyo) 138, 711-719.


[25] Sato, S., Sanjo, H., Takeda, K., Ninomiya-Tsuji, J., Yamamoto, M., Kawai, T,


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Fig. 1. PP2Cβ is covalently modified with ISG15. The extracts of HeLa cells that had been transiently transfected with indicated plasmids were subjected to Western blotting with anti-T7 tag and anti-S peptide antibodies (left panel) or to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-Flag tag and anti-T7 tag antibodies (right panel). The ISGylated PP2Cβ is indicated by an open arrowhead. Nonspecific bands are indicated by asterisks.

Fig. 2. PP2Cβ is ISGylated in response to IFN signal. (A) The extract of HeLa cells that had been transiently transfected with Flag-tagged PP2Cβ expression plasmid, followed by 24-h treatment with IFNβ (1,000 units/ml) (Pepro Tech), was subjected to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-Flag tag and anti-ISG15 antibodies. (B) The extract of HeLa cells pretreated with IFNβ was subjected to immunoprecipitation with anti-PP2Cβ antibody and control immunoglobulin (Ig) and then to Western blotting with anti-PP2Cβ and anti-ISG15 antibodies.

Fig. 3. ISGylation system suppresses PP2Cβ activity. HeLa cells were transiently transfected with indicated plasmids together with pNF-κB reporter plasmid and were then lysed, and
luciferase activity was measured. Results are expressed as fold induction in luciferase activity relative to that of control mocked cells that had been transfected with an empty vector. Triple experiments were carried out, and the error bars represent standard deviation. The ISGylation system contains T7-tagged ISG15, S-tagged UBE1L, and S-tagged UbcH8.

Fig. 4. ISGylation of PP2Cβ deletion mutants. The extracts of HeLa cells expressing the wild type PP2Cβ and its various deletion mutants, (A) Δ1-295 and Δ296-479 and (B) Δ1-92, Δ93-193, Δ194-295, Δ296-378 and Δ379-479, in the presence and absence of the ISGylation system were subjected to immunoprecipitation with anti-Flag tag antibody and then to Western blotting with anti-Flag tag and anti-T7 tag antibodies.

Fig. 5. ISGylation of PP2Cβ KR mutants. The extracts of HeLa cells expressing various KR mutants, N-terminally Flag-tagged (A) and C-terminally Flag-tagged (B), in the presence of the ISGylation system were subjected to immunoprecipitation and then to Western blotting as in Fig. 4.

Fig. 6. ISGylation of PP2Cβ suppresses its activity. (A) HeLa cells were transiently transfected with indicated plasmids together with pNF-κB reporter plasmid, and luciferase activity was measured as in Fig. 3. (B) The extracts of HeLa cells that had been transiently
transfected with indicated plasmids and then treated with cycloheximide (CHX) (50 µg/ml) for 30 min were subjected to Western blotting with anti-IκBα, anti-Flag tag, anti-S peptide, and anti-T7 tag antibodies. Wt, wild type; KR, K12,142R.
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