

SUMOylation Attenuates Human β -Arrestin 2 Inhibition of IL-1R/TRAF6 Signaling*

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Background: It was unknown whether human β -arrestin 2 could be SUMOylated.

Results: Human β -arrestin 2 is SUMOylated on Lys-295. SUMOylation attenuates β -arrestin 2 inhibition of IL-1R/TRAF6 signaling.

Conclusion: SUMOylation attenuates human β -arrestin 2 inhibition of TRAF6 and IL-1R signaling.

Significance: We show SUMOylation as a novel mechanism in regulation of β -arrestin 2-mediated IL-1R-TRAF6 signaling.

β -Arrestin 2 as an adaptor plays a role in the regulation of receptor desensitization, trafficking, and signaling. Bovine β -arrestin 2 has been shown to be SUMOylated on the lysine 400 residue, which links it to the endocytosis of the β_2 -adrenergic receptor. Here we identify a major SUMOylation site, lysine 295, on human β -arrestin 2. SUMOylation on this site attenuates β -arrestin 2 binding to TRAF6, then enhances TRAF6 oligomerization and autoubiquitination, and consequently leads to the increase of TRAF6-mediated NF- κ B/AP-1 activation. We further determine SENP1 as a specific de-SUMOylation protease that can reverse the SUMOylation of β -arrestin 2-mediated processes. Our study reveals SUMOylation as a novel mechanism in the regulation of β -arrestin 2-mediated IL-1R/TRAF6 signaling.

IL-1 signaling is a key player in the regulation of inflammatory processes. IL-1 stimulates IL-1R and downstream signaling molecules and subsequently activates the transcription factors NF- κ B and AP-1, which control the expression of key immunoregulatory genes (1–3). TRAF6 is a critical mediator for the Toll-like/interleukin-1 receptor superfamily (4, 5). As a RING domain containing E3 ubiquitin ligase, TRAF6 is recruited to the receptor complexes and forms oligomers upon signaling activation and then leads to Lys-63-linked polyubiquitination of itself and downstream signaling molecules (6–8). Lys-63 ubiquitin-conjugated TRAF6 recruits TAB2 and activates the TAB2-associated TAK1 kinase, which subsequently phosphorylates and activates I κ B kinases. I κ B kinase then phosphorylates

I κ B α , leading to degradation of I κ B α and, consequently activation of NF- κ B. In addition, TAK1 can also activate the JNK and p38 MAPK family members, then triggering AP-1 activation (8–10). Both of oligomerization and autoubiquitination are critical for TRAF6 activity toward downstream targets to mediate IL-1 β - or LPS-induced NF- κ B/AP-1 activation.

β -Arrestin 2 (also known as arrestin 3), along with β -arrestin 1 (arrestin 2) and visual rod (arrestin 1) and cone (arrestin 4) arrestins, is part of a small family of cytosolic adaptor proteins functioning as crucial adaptors/mediators in the regulation of a variety of cell surface receptor desensitization, trafficking, and signaling activities (11–13). β -Arrestins have been shown to be modified by phosphorylation, ubiquitination, and nitrosylation (14–16). These modifications have linked β -arrestins to G protein-coupled receptor (GPCR)³ internalization, trafficking and signal transduction (13). Additionally, β -arrestin 2 directly interacts with TRAF6 after TLR/IL-1R activation, leading to prevention of autoubiquitination of TRAF6 and activation of NF- κ B and AP-1 (17).

Recently, bovine β -arrestin 2 has been reported as a SUMOylated protein. The SUMOylation site Lys-400 locates within the C tail of bovine β -arrestin 2, which mediates GPCR endocytosis, and the SUMOylation of bovine β -arrestin 2 has been shown to promote β -arrestin 2-mediated β_2 -adrenergic receptor internalization (18). Here we also show human/murine β -arrestin 2 as a SUMOylated target. However, lysine 295, not lysine 400, has been identified as a major SUMO-conjugated site on human β -arrestin 2. We further found that SUMOylation decreases β -arrestin 2 inhibition of TRAF6 oligomerization and autoubiquitination and, consequently, promotes TRAF6-mediated NF- κ B/AP-1 activation. Our study reveals SUMOylation as a novel mechanism in the regulation of β -arrestin 2-mediated IL-1R/TRAF6 signaling.

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³ The abbreviations used are: GPCR, G protein-coupled receptor; SUMO, small ubiquitin-like modifier; MEF, mouse embryonic fibroblast; WB, Western blot.

SUMOylation Regulates β -Arrestin2 Activity

EXPERIMENTAL PROCEDURES

Plasmids and Antibodies—The plasmids β -arrestin 2-HA, FLAG-TRAF6, and His-ubiquitin were provided by Dr. Ping Wang (Institute of Biomedical and Science, East China Normal University, China). FLAG-SUMO1, FLAG-SEN1, HA-SUMO1, RGS-SEN1, and RGS-SEN1-mu, a SEN1 catalytic mutant, have been described previously (19–21). β -Arrestin 2 point mutants (K295R and K400R) were generated using site-directed mutagenesis. SEN1-GFP, SEN1mu-GFP, and GFP-TRAF6 were generated using standard cloning procedures and PCR-based mutagenesis (Vazyme Biotech Co., Ltd). Antibodies against FLAG M2 and HA were from Sigma; GFP from Eptomics; RGS-his from Qiagen; and β -arrestin 2, $\text{I}\kappa\text{B}\alpha$, SUMO1, ERK, and p-ERK from CST.

Immunoprecipitation and Immunoblotting—Transfected cells were lysed in radioimmune precipitation assay buffer (50 mM Tris-HCl (pH 7.4), 150 mM NaCl, 1 mM EDTA, 1% Nonidet P-40, 0.1% SDS, 1% sodium deoxycholate, and a mixture of protease inhibitors) and cleared by centrifugation. Cleared cell lysates were incubated with 10 μl of anti-FLAG M2-agarose affinity gel (Sigma) or 10 μl of anti-HA-agarose affinity gel (Sigma) for 2 h. To perform an endogenous assay, MEF cells were lysed in ice-cold radioimmune precipitation assay buffer. Cleared cell lysates were incubated with anti-SUMO-1 antibody (1:200) and 15–20 μl of protein A/G beads (Santa Cruz Biotechnology) for 4 h at 4 $^{\circ}\text{C}$. After extensive washing, beads were boiled at 100 $^{\circ}\text{C}$ for 10 min. Proteins were resolved by SDS-PAGE and transferred onto PVDF membranes (Millipore), followed by immunoblotting using corresponding antibodies according to the instructions of the manufacturer. Immunoblots were analyzed using the LAS-4000 system (Fujifilm).

Ubiquitination Assay—HEK293T cells were transfected with His-ubiquitin and FLAG-TRAF6. The transfected cells were lysed by denatured buffer (6 M guanidine-HCl, 0.1 M $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$, and 10 mM imidazole), followed by Talon bead purification. The ubiquitination was detected by Western blot analysis.

Luciferase Assay—NF- κB - or AP-1-dependent firefly luciferase plasmids were transiently transfected into HEK293T cells with *Renilla* luciferase plasmids and others. The cells were harvested 36 h after transfection, and luciferase assays were performed using the Dual-Luciferase reporter assay system (Promega). The relative luciferase activity was normalized on the basis of the *Renilla* luciferase activity. Data represent three independent experiments performed in duplicate.

Stable Cell Line— β -Arrestin 2 WT or β -arrestin 2 K295R mutant (K295R) lentiviral plasmids were transfected into HEK293T cells with lentivirus packaging vectors by calcium phosphate-DNA coprecipitation method. Viral supernatants were collected 48 h after transfection. MCF-7 cells were infected by lentiviral supernatant in the presence of 10 $\mu\text{g}/\text{ml}$ Polybrene for 12 h. 48–72 h later, the cells were sorted for stable cell lines by flow cytometry.

RNA Isolation and Real-time RT-PCR—Total RNA was isolated from cells by using Tripure isolation reagent (Roche). For mRNA analysis, an aliquot containing 2 μg of total RNA was reverse-transcribed using the cDNA synthesis kit (Takara).

Real-time PCR was performed using SYBR Green PCR master mix (Applied Biosystems) and detected by the ABI Prism 7500 sequence detection system (Applied Biosystems). The primers for real-time RT-PCR were as follows: GAPDH, 5'-GAGCTG-AACGGGAAGCTCACTG-3' (sense) and 5'-TGGTGCTCAGTGATAGCCAGGA-3' (antisense); TNF α , 5'-CCCTCTGG-CCCAGGCAGTCA-3' (sense) and 5'-ATGGGTGGAGGGG-CAGCCTT-3' (antisense).

ELISA Assay—After serum starvation for 12 h, MCF-7 cells were cultured for 24 h with recombinant human IL-1 β (20 ng/ml) (Bioworld Technology). The concentration of TNF α in culture supernatants was determined with a human-specific ELISA kit (ExCell Bio), followed by analysis with a SYNERGY microplate reader (BioTek).

RESULTS

SUMO Conjugates Human β -Arrestin 2 on Lys-295—Wyatt *et al.* (18) have reported previously that bovine β -arrestin 2 is conjugated by SUMO on residue Lys-400. However, when aligning the bovine, human, and murine β -arrestin 2 sequences, we noticed that the Lys-400 residue in human/murine β -arrestin 2 is not in a conserved SUMO consensus motif, ΨKXE (Fig. 1A), which raised the question of whether human/murine as well as bovine β -arrestin 2 could be SUMOylated. To test this, we performed an *in vivo* SUMOylation assay in HEK293T cells by cotransfecting human β -arrestin 2-HA and FLAG-SUMO1. As shown in Fig. 1, B and C, the SUMOylated β -arrestin 2 band was readily detected in cells transfected with β -arrestin 2-HA and FLAG-SUMO1. We also observed that endogenous human/mouse β -arrestin 2 was modified by endogenous SUMO1 in HEK293T cells and MEFs (Fig. 1D and Fig. 3C). These results showed that human/murine β -arrestin 2 could be also conjugated by SUMO similarly as bovine β -arrestin 2.

We mutated Lys-400 on human β -arrestin 2 to test whether this residue was a SUMO conjugation site, as bovine β -arrestin 2. As shown in Fig. 1E, mutating the Lys-400 residue had no significant effect on human β -arrestin 2 SUMOylation. However, when mutating Lys-295, a lysine residue located in a conserved SUMO consensus motif, LKHE (ΨKXE), human β -arrestin 2 SUMOylation was significantly abolished, suggesting that Lys-295, but not Lys-400, on human β -arrestin 2 is the major residue to accept SUMO.

SUMOylation Attenuates Human β -Arrestin 2 Inhibition of TRAF6 Activation—Lys-400 resides within the C tail of bovine β -arrestin 2, which mediates GPCR endocytosis. SUMOylation on this site has been shown to promote β -arrestin 2-mediated β_2 -adrenergic receptor internalization (18). However, Lys-295 on human β -arrestin 2 locates in the TRAF6-binding domain. β -Arrestin 2 binding to TRAF6 has been shown to inhibit autoubiquitination of TRAF6 and activation of NF- κB and AP-1 responding to TLR-IL-1R signaling (17). We therefore proposed that SUMOylation on Lys-295 might regulate β -arrestin 2 interacting with TRAF6 and then modulate IL-1R/TRAF6 signaling. We first tested whether β -arrestin 2 SUMOylation is related to IL-1 β stimulation. We showed that IL-1 β induced human β -arrestin 2 SUMOylation (Fig. 2A). We then determined whether SUMOylation of human β -arrestin 2 could affect TRAF6 ubiquitination. As shown in Fig. 2B, the

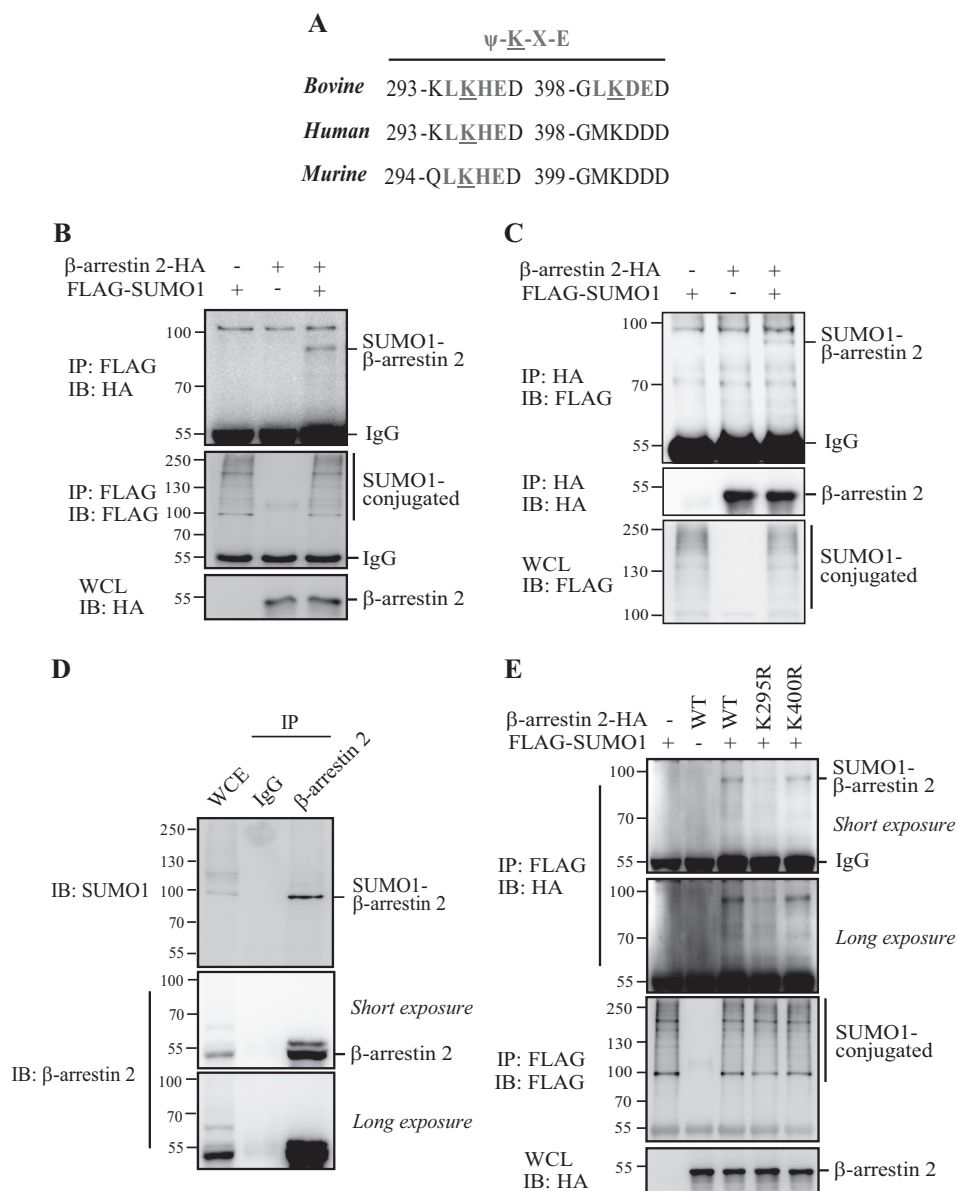


FIGURE 1. SUMO conjugates human β -arrestin 2 on Lys-295. *A*, the alignment of bovine/human/murine β -arrestin 2 sequences sowing the SUMO consensus motif. *B* and *C*, human β -arrestin 2 is SUMOylated *in vivo*. HEK293T cells were transfected with the indicated plasmids, and the transfected cells were harvested 36 h after transfection and immunoprecipitated with anti-FLAG M2-agarose beads or anti-HA beads. The immunoprecipitates (*IP*) and the original whole-cell lysates (*WCL*) were analyzed by immunoblotting (*IB*) with anti-HA or anti-FLAG antibodies. *D*, HEK293T cells lysates were immunoprecipitated with anti- β -arrestin 2 antibody or control IgG. The immunoprecipitates and the whole-cell lysates were analyzed by immunoblotting with anti-SUMO1 or anti- β -arrestin 2 antibodies. *E*, Lys-295 is the major SUMOylation site of human β -arrestin 2. HEK293T cells were transfected with the indicated plasmids, and the transfected cell lysates were immunoprecipitated with anti-FLAG M2-agarose beads, followed by Western blot analysis by using anti-HA or anti-FLAG antibodies.

expression of β -arrestin 2 markedly reduced TRAF6 ubiquitination (*lane 3 versus lane 2*). Interestingly, the expression of SUMO1/Ubc9 could almost restore TRAF6 ubiquitination (Fig. 2*B*, *lane 4 versus lane 3*) but had no effect on β -arrestin 2 K295R-mutant (Fig. 2*B*, *lane 5 versus lane 4*). Mutating Lys-400 on human β -arrestin 2 did not affect TRAF6 ubiquitination (Fig. 2*B*, *lane 6 versus lanes 3 and 4*), suggesting that SUMOylation on Lys-295 reduces β -arrestin 2 inhibition of TRAF6 ubiquitination. We further tested whether SUMOylation of β -arrestin 2 had a similar effect on TRAF6-mediated NF- κ B and AP-1 activation as on TRAF6 ubiquitination. HEK293T cells were transfected with an NF- κ B- or AP-1-dependent luciferase reporter plus TRAF6 and with or without SUMO1/Ubc9. A luciferase assay

from these transfected cells showed that SUMOylation on Lys-295 also attenuates β -arrestin 2 inhibition of NF- κ B and AP-1 activation (Fig. 2*C*).

We generated human β -arrestin 2 wild-type, K295R-mutant, or K400R-mutant stably transfected MCF-7, a relatively low endogenous β -arrestin 2 cell line (data not shown), to further determine the role of SUMOylation in β -arrestin 2-mediated IL-1R signaling. As shown in Fig. 2*D*, IL-1 β -induced TRAF6 ubiquitination was significantly lower in β -arrestin 2 (K295R)-MCF-7 cells than in β -arrestin 2 (WT)-MCF-7 or β -arrestin 2 (K400R)-MCF-7 cells. Similarly, IL-1 β -induced NF- κ B or AP-1 activation was also significantly lower in β -arrestin 2 (K295R)-MCF-7 cells than that in β -arrestin 2 (WT)-MCF-7 or β -arres-

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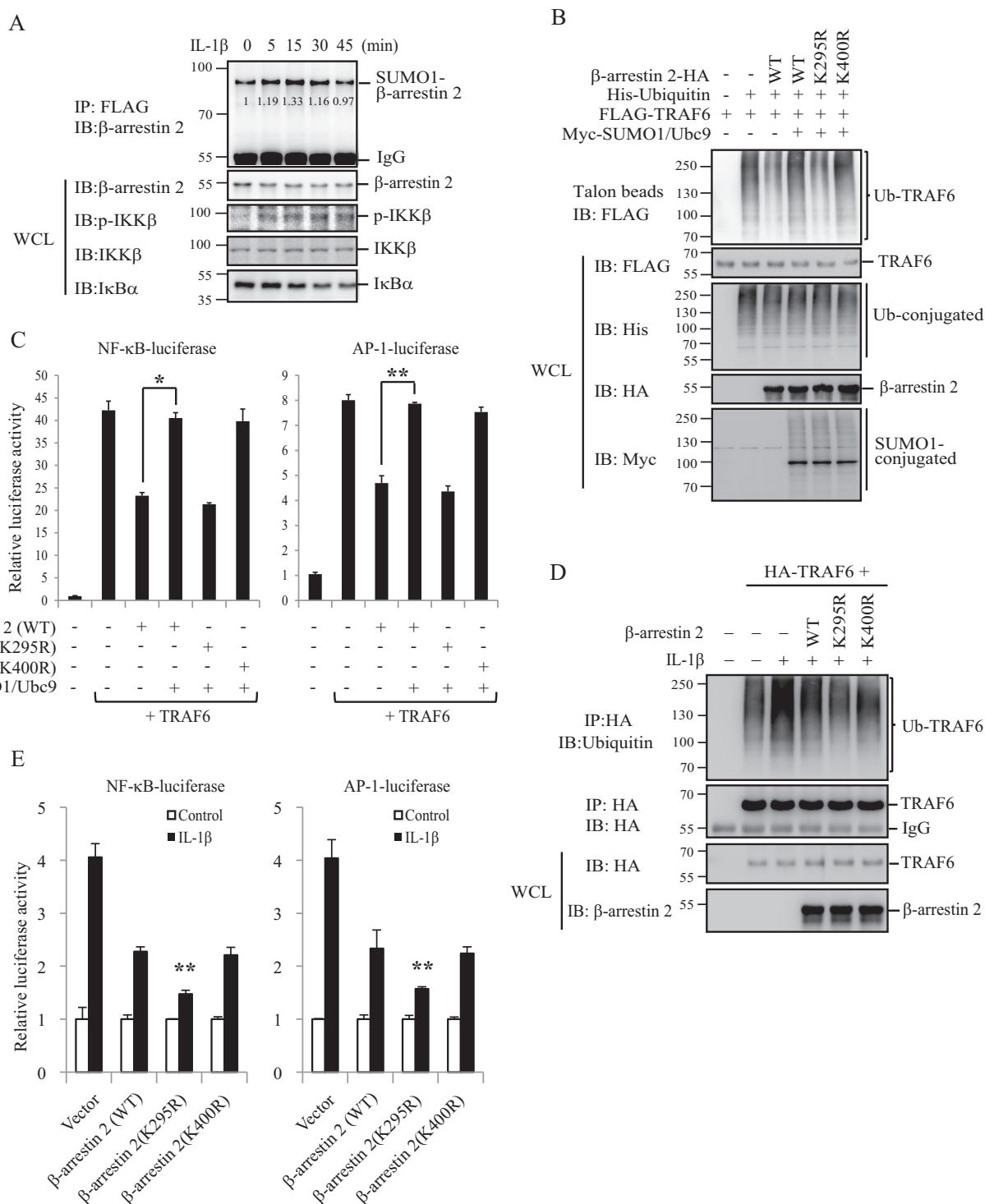


FIGURE 2. SUMOylation attenuates human β -arrestin 2 inhibition of TRAF6 activation. *A*, SUMOylation of β -arrestin 2 is induced by IL-1 β . HEK293T cells were transfected with FLAG-SUMO1 and β -arrestin 2-HA. The transfected cells were treated with IL-1 β (20 ng/ml) for the indicated time before harvest. The cell lysates were immunoprecipitated (IP) with anti-FLAG M2-agarose beads and analyzed by immunoblotting (IB) with anti- β -arrestin 2, anti-p-IKK β , anti-IKK β , or anti-I κ B α antibodies. IKK β phosphorylation and I κ B α degradation were used as indicators for the validity of IL-1 β stimuli. The relative gray scale determination was analyzed using ImageJ software. *WCL*, whole-cell lysate. *B*, SUMOylation reduces β -arrestin 2 inhibition of TRAF6 autoubiquitination. The indicated plasmids were transfected into HEK293T cells. The Talon bead precipitates and cell lysates were analyzed by immunoblotting with anti-FLAG, anti-HA, anti-His, or anti-Myc antibodies. *Ub*, ubiquitin. *C*, SUMOylation attenuates β -arrestin 2 inhibition of TRAF6-mediated NF- κ B and AP-1 activation. An NF- κ B- or AP-1-dependent firefly luciferase reporter and a *Renilla* luciferase reporter plus the indicated plasmids were cotransfected into HEK293T cells. The relative luciferase activity was measured 36 h after transfection and normalized on the basis of *Renilla* luciferase activity. *, $p < 0.05$; **, $p < 0.01$. *D*, β -arrestin 2 WT-, β -arrestin 2 K295R-, or β -arrestin 2 K400R-MCF-7 stable cells were transfected with HA-TRAF6 or not as indicated. The cells were treated with IL-1 β . The cell lysates were immunoprecipitated with anti-HA-agarose beads, and the immunoprecipitates and the cell lysates were analyzed by immunoblotting with anti-ubiquitin, anti-HA, or anti- β -arrestin 2 antibodies. *E*, β -arrestin 2 WT-, β -arrestin 2 K295R-, or β -arrestin 2 K400R-MCF-7 stable cells were transfected with an NF- κ B- or AP-1-dependent firefly luciferase reporter and a *Renilla* luciferase reporter. The relative luciferase activity was measured 36 h after transfection and normalized on the basis of *Renilla* luciferase activity. *, $p < 0.05$; **, $p < 0.01$.

tin 2 (K400R)-MCF-7 cells (Fig. 2E). These results suggest that SUMOylation on Lys-295 could attenuate β -arrestin 2 inhibition of TRAF6 activation and TRAF6-mediated IL-1R signaling.

SEN1 De-SUMOylates β -Arrestin 2—We observed that SUMO-specific protease 1 (SEN1) could bind to β -arrestin 2 (Fig. 3A). Therefore, we reasoned that SEN1 could be a deconjugation protease for SUMOylated β -arrestin 2. To test this possibility, we performed a de-SUMOylation assay in the transfected cells. As shown in Fig. 3B, the expression of SEN1 wild-type but not the SEN1 catalytic mutant made the SUMOylated β -arrestin 2 band disappear. We further confirmed SEN1 as a specific de-SUMOylation protease of β -arrestin 2 by observing the accumulation of the SUMOylated β -arrestin 2 in *Senp1*^{-/-} MEFs (Fig. 3C).

Because SUMOylation reduces β -arrestin 2 inhibition of TRAF6 autoubiquitination and TRAF6-mediated NF- κ B/AP-1 activation (Fig. 2), we speculated that SEN1 could enhance human β -arrestin 2 inhibition of TRAF6 ubiquitination and activation through de-SUMOylation. To test this, we performed an ubiquitination assay in HEK293T cells that were cotransfected with FLAG-TRAF6 and His-Ubiquitin plus β -arrestin 2-HA in the presence of SEN1-GFP wild-type or the catalytic mutant SEN1mu-GFP. As shown in Fig. 3D, the coexpression of SEN1 wild-type, not the catalytic mutant, and β -arrestin 2 decreased TRAF6 ubiquitination more than expression of β -arrestin 2 alone. Interestingly, SEN1-reduced TRAF6 ubiquitination depended on β -arrestin 2 because SEN1 action on TRAF6 ubiquitination could not be detected without the coexpression of β -arrestin 2. We also assessed the effect of SEN1 on TRAF6-mediated NF- κ B and AP-1 activation by using an NF- κ B- or AP-1-dependent luciferase assay. Similarly, SEN1 wild-type, not the catalytic mutant, can enhance β -arrestin 2 inhibition of TRAF6-mediated NF- κ B or AP-1 activation (Fig. 3E).

We further determined the role of SEN1 in IL-1R signaling by using *Senp1*^{-/-} MEFs. As shown in Fig. 3F, IL-1 β induced more TRAF6 ubiquitination in *Senp1*^{-/-} MEFs than that in wild-type MEFs. We also observed that IL-1 β -induced more TRAF6 ubiquitination in SEN1 knockdown β -arrestin 2 stably transfected MCF-7 cells (Fig. 3G). More interestingly, there was no effect on β -arrestin 2 (K295R)-MCF-7 cells (Fig. 3G, lane 2 versus lane 4), indicating that SEN1 negative regulation of TRAF6 ubiquitination is dependent on de-SUMOylation of β -arrestin 2. Similarly, IL-1 β induced more I κ B α phosphorylation and ERK phosphorylation in *Senp1*^{-/-} MEFs than that in wild-type MEFs (Fig. 3H). These data suggest that SEN1 enhances β -arrestin 2 inhibition of TRAF6-mediated signaling through de-SUMOylation of β -arrestin 2.

SUMOylation Decreases β -Arrestin 2 Binding to TRAF6— β -Arrestin 2 has been reported to directly interact with TRAF6 after TLR-IL-1R activation, leading to prevention of TRAF6 oligomerization and autoubiquitination (17). Because SUMOylation on Lys-295 attenuates β -arrestin 2 inhibition of TRAF6 autoubiquitination and activation (Fig. 2), we reasoned that SUMOylation might disrupt β -arrestin 2-TRAF6 interaction, which reduces β -arrestin 2 inhibition of TRAF6. To test this, we showed that the coexpression of SUMO1/Ubc9, which pro-

motes β -arrestin 2 SUMOylation, could decrease both β -arrestin 2 wild-type and β -arrestin 2 K400R-mutant binding to TRAF6 (Fig. 4A, lanes 3 and 5 versus lane 2) but has no effect on β -arrestin 2 K295R-mutant (Fig. 4A, lane 4 versus lane 2). Furthermore, we assessed β -arrestin 2-TRAF6 interaction in wild-type and *Senp1*^{-/-} MEFs. As shown in Fig. 4B, TRAF6 was able to bind to β -arrestin 2 in *Senp1*^{-/-} MEFs much less than that in wild-type MEFs. These results suggest that SUMOylation on Lys-295 could attenuate β -arrestin 2 binding to TRAF6.

Because oligomerization is necessary for TRAF6 autoubiquitination (8), we reasoned that SUMOylation of β -arrestin 2 would enhance TRAF6 oligomerization by attenuating β -arrestin 2 interaction with TRAF6. To test this, we cotransfected differently tagged TRAF6 into HEK293T cells. The coexpression of β -arrestin 2 reduced the association of GFP-TRAF6 with FLAG-TRAF6 (Fig. 4C, lane 3 versus lane 2). Interestingly, the β -arrestin 2 K295R-mutant blocked TRAF6 oligomerization more efficiently than β -arrestin 2 wild-type did (Fig. 4C, lane 4 versus lane 3), suggesting that SUMOylation of β -arrestin 2 could enhance TRAF6 oligomerization through disrupting the interaction between β -arrestin 2 and TRAF6.

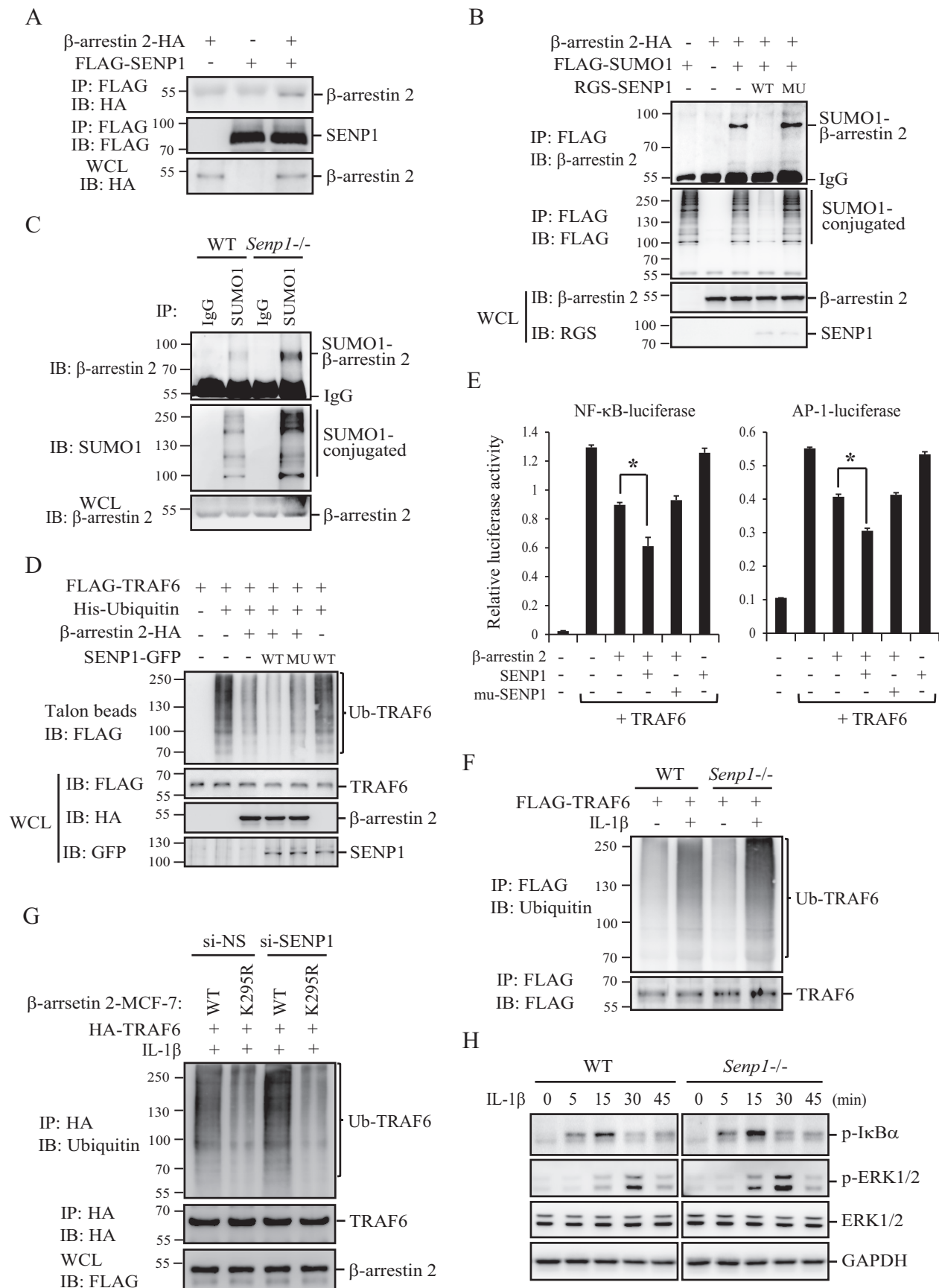
Deficiency in SUMOylation Enhances β -Arrestin 2 Inhibition of IL-1 β -induced TNF α Expression—To determine the significance of β -arrestin 2 SUMOylation *in vivo*, we compared TNF α expression in β -arrestin 2 (WT)-MCF7 and β -arrestin 2 (K295R)-MCF7 cells. As shown in Fig. 5A, the expression of β -arrestin 2 wild-type reduced IL-1 β -induced TNF α mRNA expression. However, the expression of β -arrestin 2 K295R-mutant showed more of a reduction of the TNF α mRNA level than β -arrestin 2 wild-type. Furthermore, the ELISA assay confirmed that IL-1 β -treated β -arrestin 2 (K295R)-MCF7 cells produced much less TNF α in culture medium than β -arrestin 2 (WT)-MCF7 cells (Fig. 5B). Taken together, these results indicate that SUMOylation attenuates β -arrestin 2 inhibition of IL-1 β -induced TNF α expression.

DISCUSSION

In this study, we identify Lys-295 as a major SUMOylation site on human β -arrestin 2. Because this site locates in the TRAF6-binding domain, SUMOylation can attenuate β -arrestin 2 binding to TRAF6. Therefore, SUMOylation of β -arrestin 2 can enhance TRAF6 oligomerization and autoubiquitination and, consequently, activate TRAF6-mediated IL-1R signaling. We also found SEN1 as a specific de-SUMOylation protease of β -arrestin 2 in these processes to enhance β -arrestin 2 inhibition of TRAF6 activation. These data reveal SUMOylation as a novel mechanism to attenuate β -arrestin 2 inhibition of TRAF6 and IL-1R signaling.

As an important adaptor, β -arrestins are widely involved in receptors, especially GPCR family, desensitization, trafficking, signaling, and regulating a growing list of cellular processes such as chemotaxis, apoptosis, inflammatory processes, and metastasis (13, 22). More recently, studies have shown that β -arrestins function as scaffold proteins for many signaling molecules in the cytoplasm and nucleus, thereby regulating gene expression and cellular responses (23). Besides traditional GPCR signaling, β -arrestins participate in the regulation of other signaling pathways, including TLR/IL-1R signaling and

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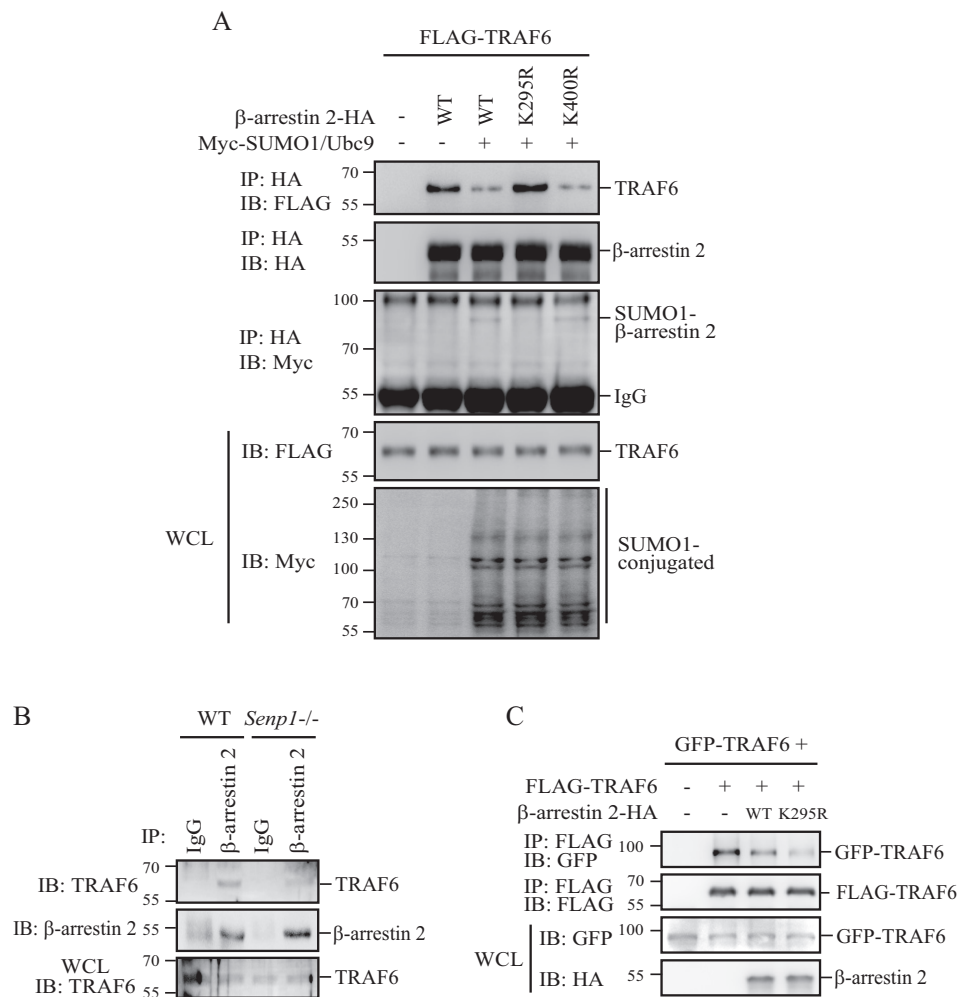


FIGURE 4. SUMOylation decreases β -arrestin 2 binding to TRAF6. A, HEK293T cells were transfected with the indicated plasmids. The cell lysates were immunoprecipitated (IP) with anti-HA-agarose beads. The immunoprecipitates and cell lysates were analyzed by WB with anti-FLAG, anti-HA, or anti-Myc antibodies. IB, immunoblotting; WCL, whole-cell lysate. B, *Senp1*^{-/-} MEFs or wild-type MEF cell lysates were immunoprecipitated with anti- β -arrestin 2 antibody or control IgG. Precipitates and cell lysates were analyzed by WB with anti- β -arrestin 2 or anti-TRAF6 antibodies. C, GFP-TRAF6 and FLAG-TRAF6 were cotransfected into HEK293T cells with β -arrestin 2-HA wild-type or K295R-mutant, and the transfected cells were harvested 24 h after transfection. The cell lysates were immunoprecipitated with anti-FLAG M2-agarose beads. The precipitates and cell lysates were analyzed by WB with anti-GFP, anti-FLAG, or anti-HA antibodies.

Wnt/ β -catenin signaling (17, 24–26). In TLR/IL-1R signaling, β -arrestin 2 has been reported to directly bind to TRAF6 and then reduce TRAF6 autoubiquitination and activation, leading to a negative regulation of TLR/IL-1R signaling. Here we further find β -arrestin 2 SUMOylation, as a negative mechanism,

to attenuate β -arrestin 2 inhibition of TRAF6 activation and then promote TRAF6 oligomerization and autoubiquitination.

SUMOylation is an essential posttranslational modification with critical roles in regulating protein functions such as localization, activity, and protein-protein interaction (27–29). Typ-

FIGURE 3. SENP1 de-SUMOylates β -arrestin 2. A, SENP1 binds to β -arrestin 2. HEK293T cells were transfected with the indicated plasmids, and the transfected cell lysates were immunoprecipitated (IP) with anti-FLAG M2-agarose beads, followed by WB analysis with anti-HA or anti-FLAG antibodies. IB, immunoblot; WCL, whole-cell lysate. B, SENP1 deconjugates SUMOylated β -arrestin 2 *in vivo*. HEK293T cells were transfected with the indicated plasmids, and the transfected cell lysates were immunoprecipitated with anti-FLAG M2-agarose beads. The immunoprecipitates and the whole-cell lysates were analyzed by immunoblotting with anti- β -arrestin 2, anti-FLAG, or anti-RGS-his antibodies. MU, mutant. C, SUMOylated β -arrestin 2 is accumulated in *Senp1*^{-/-} MEFs. MEF cell lysates were immunoprecipitated with anti-SUMO1 antibody or control IgG. The immunoprecipitates and cell lysates were analyzed by immunoblotting with anti- β -arrestin 2 or anti-SUMO1 antibodies. D, SENP1 enhances β -arrestin 2 inhibition of TRAF6 autoubiquitination. The indicated plasmids were transfected into HEK293T cells. The cell lysates were purified by Talon beads. The precipitates and cell lysates were immunoblotted using anti-FLAG, anti-HA, or anti-GFP antibodies. E, SENP1 enhances β -arrestin 2 inhibition of TRAF6-mediated NF- κ B and AP-1 activation. The indicated plasmids were cotransfected into HEK293T cells with an NF- κ B- or AP-1-dependent firefly luciferase reporter and a *Renilla* luciferase reporter. The relative luciferase activity was measured 36 h after transfection and normalized on the basis of *Renilla* luciferase activity. *, $p < 0.05$; **, $p < 0.01$. F, FLAG-TRAF6-transfected wild-type and *Senp1*^{-/-} MEFs were treated with IL-1 β or left untreated before harvest. The cell lysates were immunoprecipitated with anti-FLAG M2-agarose beads, followed by WB analysis with anti-ubiquitin or anti-TRAF6 antibodies. G, β -arrestin 2 WT- or β -arrestin 2 K295R-MCF-7 stable cells were transfected with HA-TRAF6 and si-SENP1 or si-NS oligonucleotides. The cells were treated with IL-1 β before harvest, the cell lysates were immunoprecipitated with anti-HA-agarose beads, and the immunoprecipitates and the cell lysates were analyzed by immunoblotting with anti-ubiquitin, anti-HA, or anti-FLAG antibodies. H, wild-type and *Senp1*^{-/-} MEFs were treated with IL-1 β for the indicated time before harvest, followed by WB analysis with the indicated antibodies.

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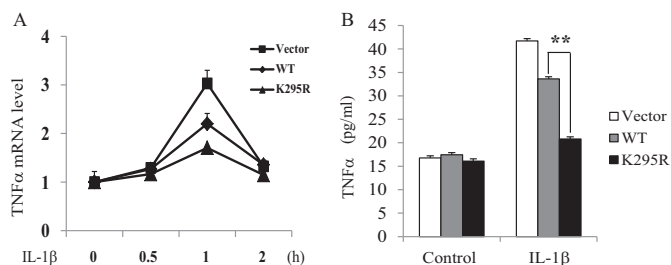


FIGURE 5. Deficiency in SUMOylation enhances β -arrestin 2 inhibition of IL-1 β -induced TNF α expression. *A*, quantitative real-time RT-PCR analysis showing the relative TNF α mRNA level in β -arrestin 2 WT- or β -arrestin 2 K295R-MCF-7 stable cell lines treated with IL-1 β for the indicated time. The results are representative of three experiments. *B*, ELISA analysis showing the TNF α concentration in supernatants of β -arrestin 2 WT- or β -arrestin 2 K295R-MCF-7 stable cell lines after IL-1 β stimulation. The results are representative of three experiments. **, $p < 0.01$.

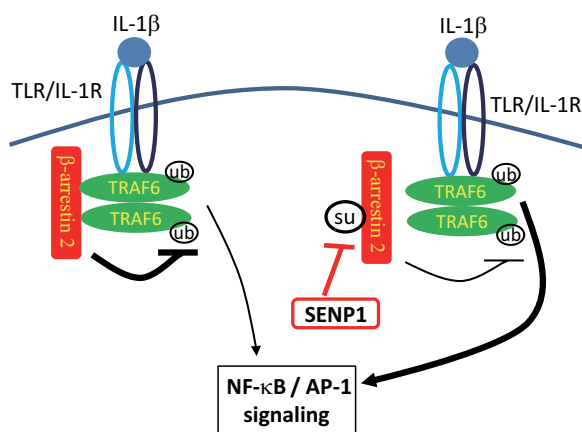


FIGURE 6. Working model for the regulation of IL-1R-TRAF6 signaling by β -arrestin 2 SUMOylation.

ically, SUMO-modified proteins contain a SUMO consensus motif defined as Ψ KXE, where Ψ is a large hydrophobic residue and X represents any amino acid (30). Bovine β -arrestin 2 has been shown to be SUMOylated. SUMOylation of bovine β -arrestin 2 does not influence the affinity of receptor interaction but is important for AP-2 interaction and AP-2-mediated receptor internalization (18). However, the Lys-400 residue in the carboxyl-terminal region of human β -arrestin 2 is not a major SUMOylation site because mutation of this site did not significantly affect its SUMOylation status. We identified Lys-295 as a major SUMOylation site on human β -arrestin 2, although there might be other potential SUMOylation sites. This site locates in the TRAF6-binding domain of human β -arrestin 2. Therefore, we observed that SUMOylation of human β -arrestin 2 plays a role in the regulation of TRAF6 activation and TRAF6-mediated signaling. It would be interesting to study whether it is evolution-related that the different SUMOylation sites exist on bovine and human/murine β -arrestin 2. In summary, our study reveals that SUMOylation modulates human β -arrestin 2-mediated inhibition of TRAF6 and TRAF6-mediated IL-1R signaling (Fig. 6).

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