

Lysine 63-linked Polyubiquitination of TAK1 at Lysine 158 Is Required for Tumor Necrosis Factor α - and Interleukin-1 β -induced IKK/NF- κ B and JNK/AP-1 Activation*[§]

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Transforming growth factor- β -activated kinase 1 (TAK1) plays an essential role in the tumor necrosis factor α (TNF α)- and interleukin-1 β (IL-1 β)-induced I κ B kinase (IKK)/nuclear factor- κ B (NF- κ B) and c-Jun N-terminal kinase (JNK)/activator protein 1 (AP-1) activation. Here we report that TNF α and IL-1 β induce Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue within the kinase domain. Tumor necrosis factor receptor-associated factors 2 and 6 (TRAF2 and -6) act as the ubiquitin E3 ligases to mediate Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue *in vivo* and *in vitro*. Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue is required for TAK1-mediated IKK complex recruitment. Reconstitution of TAK1-deficient mouse embryo fibroblast cells with TAK1 wild type or a TAK1 mutant containing a K158R mutation revealed the importance of this site in TNF α and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation as well as IL-6 gene expression. Our findings demonstrate that Lys⁶³-linked polyubiquitination of TAK1 at Lys¹⁵⁸ is essential for its own kinase activation and its ability to mediate its downstream signal transduction pathways in response to TNF α and IL-1 β stimulation.

Tumor necrosis factor α (TNF α)³ and interleukin-1 β (IL-1 β) are two potent proinflammatory cytokines that play important roles in the regulation of immunity, inflammation, cell proliferation, differentiation, and apoptosis (1, 2). Cellular responses to

TNF α and IL-1 β are mediated by intracellular signaling pathways that control the activation of nuclear factor- κ B (NF- κ B) and activator protein 1 (AP-1) (3, 4).

Upon binding to its receptor, TNF α induces formation of a receptor-associated complex, including the adaptor proteins TRADD, TRAF2, TRAF5, and RIP1, which subsequently leads to Lys⁶³-linked polyubiquitination of TRAF2 and RIP1 (5–7). In contrast, IL-1 β binding to its receptor induces a receptor-associated complex formation, including MyD88, IRAK1, IRAK4, and TRAF6, which is followed by Lys⁶³-linked polyubiquitination of TRAF6 and IRAKs (8–12). The formation of TRAF2-RIP1 and TRAF6-IRAK4 complexes as well as the Lys⁶³-linked polyubiquitination of RIP1 and TRAF6 appear to enable the recruitment and activation of transforming growth factor- β -activated kinase 1 (TAK1) through binding of the TAK1 regulatory subunits TAB2 and TAB3 to the Lys⁶³-polyubiquitinated RIP1 and TRAF6. The activated TAK1 then triggers the activation of the I κ B kinase (IKK), c-Jun N-terminal kinase (JNK), and p38 MAPK (8, 13–17), which leads to activation of transcription factors NF- κ B and AP-1 and up-regulation of many genes encoding proinflammatory cytokines, chemokines, adhesion molecules, and proteolytic enzymes (18).

The IKK complex consists of three subunits: two catalytic subunits, IKK α and IKK β , and an essential regulatory subunit, IKK γ /NF- κ B essential modulator (NEMO) (3, 19). Genetic studies have implicated that IKK β and IKK γ /NEMO are essential for the TNF α - and IL-1 β -mediated NF- κ B activation (20–22). Phosphorylation of serine 177 and 181 residues in the activation loop is required for IKK β activation (23). IKK γ /NEMO has been indicated to bind Lys⁶³-linked polyubiquitin chains (7, 16, 24, 25). It is proposed that Lys⁶³-polyubiquitin chains act as a scaffold to allow for assembly of a signaling complex that leads to IKK β activation. Once activated, IKK β phosphorylates I κ B proteins and leads to I κ B polyubiquitination with a Lys⁴⁸-linked ubiquitin chain. Polyubiquitination-mediated degradation of I κ Bs allows NF- κ B to translocate into the nucleus and activate NF- κ B-dependent gene expression (26).

JNKs are members of three related mitogen-activated protein kinases (MAPKs), including the extracellular signal-regulated kinases (ERKs), JNKs, and p38 MAPKs (4, 27). JNKs and p38 MAPKs are involved in transmitting intracellular signals in

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³ The abbreviations used are: TNF α , tumor necrosis factor- α ; NF- κ B, nuclear factor- κ B; IKK, I κ B kinase; IL, interleukin; NEMO, NF- κ B essential modulator; JNK, c-Jun N-terminal kinase; MAPK, mitogen-activated protein kinase; TGF- β , transforming growth factor- β ; GFP, green fluorescent protein; GST, glutathione S-transferase; Ub, ubiquitin; MEF, mouse embryonic fibroblast.

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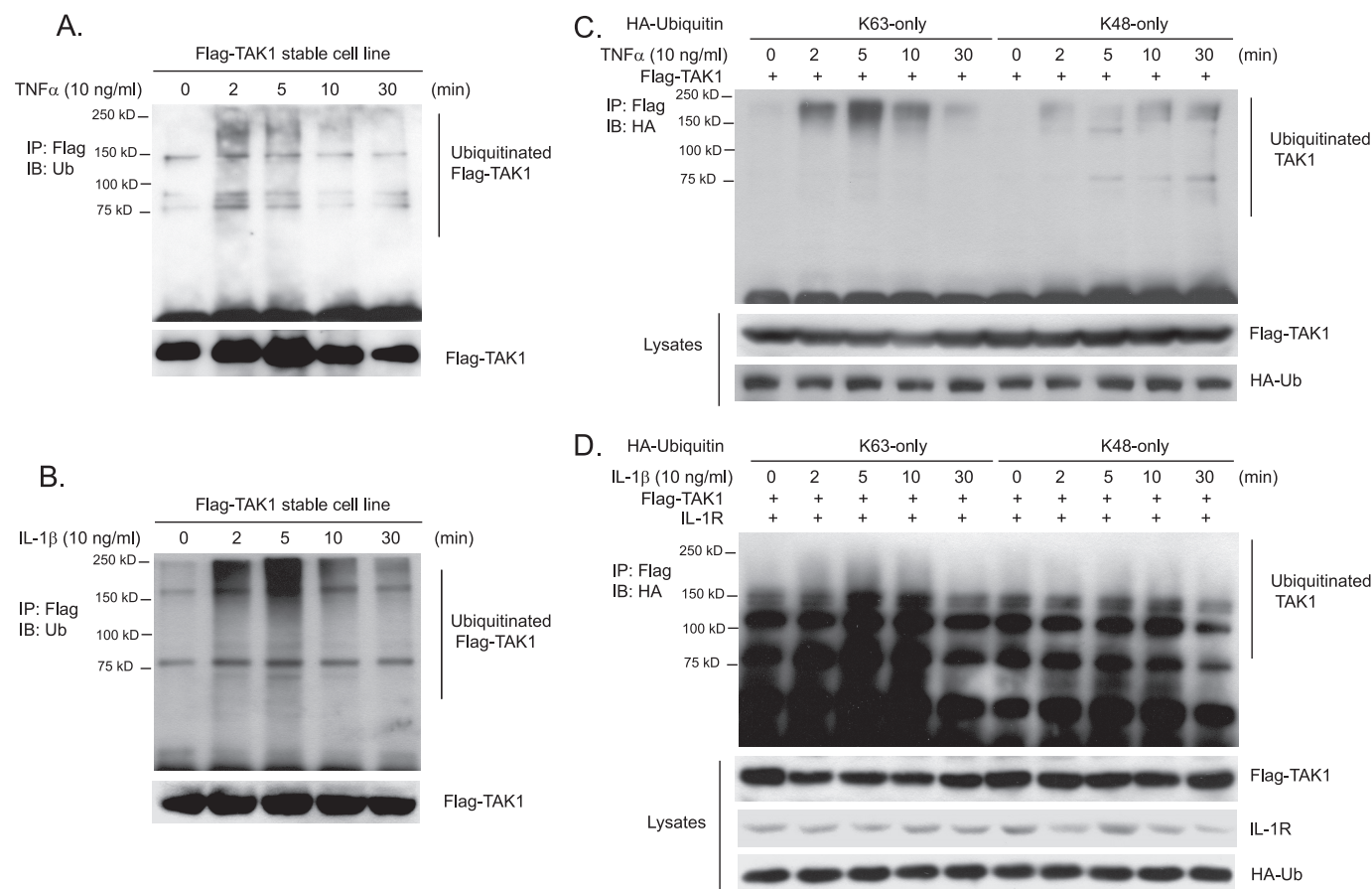


FIGURE 1. TNF α and IL-1 β induce Lys⁶³-linked TAK1 polyubiquitination. *A* and *B*, TNF α and IL-1 β induce TAK1 polyubiquitination. HeLa cells with stable expression of FLAG-TAK1 were either untreated or treated with TNF α (10 ng/ml) (*A*) and IL-1 β (10 ng/ml) (*B*) for the times indicated and subsequently lysed. FLAG-TAK1 proteins in the cell lysates were immunoprecipitated (*IP*) with anti-FLAG antibodies and immunoblotted (*IB*) with anti-ubiquitin antibodies to detect the presence of ubiquitinated FLAG-TAK1. *C*, TNF α induces Lys⁶³-linked TAK1 polyubiquitination. Expression vectors encoding FLAG-TAK1 were co-transfected into HEK-293T cells with expression vectors encoding HA-ubiquitin-Lys⁶³ only and Lys⁴⁸ only, respectively. Then cells were either untreated or treated with TNF α (10 ng/ml) for the time points indicated and subsequently lysed. FLAG-TAK1 proteins in the cell lysates were immunoprecipitated with anti-FLAG antibodies and immunoblotted with anti-HA antibodies to detect the presence of ubiquitinated FLAG-TAK1. *D*, IL-1 β induces Lys⁶³-linked TAK1 polyubiquitination. Expression vectors encoding FLAG-TAK1 and IL-1R were co-transfected into HEK-293T cells with expression vectors encoding HA-ubiquitin-Lys⁶³-only and Lys⁴⁸-only, respectively. Then cells were either untreated or treated with IL-1 β (10 ng/ml) for the times indicated and subsequently lysed. FLAG-TAK1 proteins in the cell lysates were immunoprecipitated with anti-FLAG antibodies and immunoblotted with anti-HA antibodies to detect the presence of ubiquitinated FLAG-TAK1.

response to proinflammatory cytokines, such as TNF α and IL-1 β , and environmental stresses (28). Once activated, JNKs phosphorylate specific sites on the N-terminal transactivation domain of transcription factor c-Jun, an important component of transcriptional activator AP-1. Phosphorylation of these sites stimulates the transactivity of c-Jun to activate AP-1-dependent gene expression (29).

TAK1, a member of the MAPK kinase kinase family, was originally found to function in the transforming growth factor- β (TGF- β)-mediated MAPK activation (30). TAK1 has been demonstrated to be essential in TNF α - and IL-1 β -mediated activation of NF- κ B, JNK, and p38 (15, 31–33). Several regulatory subunits of TAK1, including TAB1, TAB2, TAB3, and TAB4, have been implicated to play a role in the regulation of TAK1 activity in response to TNF α and IL-1 β stimulation (13, 34–37). TAB1 is a TAK1-interacting protein and induces TAK1 kinase activity through promoting autophosphorylation of key serine/threonine sites of the kinase activation loop (36). TAB1 is an inactive pseudophosphatase sharing homology with members of the PPM family of protein serine/threonine phos-

phatases (38). TAB2, TAB3, and TAB4 activate TAK1 through binding to polyubiquitinated proteins and promoting a larger complex formation during TNF α - and IL-1 β -induced TAK1 activation (15, 37).

Currently, growing evidence suggests that protein phosphorylation and ubiquitination play an essential role in the regulation of TAK1 activation. X-linked inhibitor of apoptosis inhibits JNK1 activation by TGF- β 1 through ubiquitin-mediated proteosomal degradation of TAK1 (39). Furthermore, several reports suggest TAK1 ubiquitination is involved in the regulation of TAK1-mediated signaling pathways (40–42). Recently, TRAF6-mediated Lys⁶³-linked TAK1 polyubiquitination at the Lys³⁴ residue has been shown to correlate with TAK1 activation in TGF- β signaling (43). *Helicobacter pylori* CagA activates NF- κ B by targeting TAK1 for TRAF6-mediated Lys⁶³-linked ubiquitination (44). However, the molecular regulation of TAK1 activation by TNF α - and IL-1 β -induced protein ubiquitination remains poorly understood.

In this report, we have further investigated the molecular mechanism of TAK1 activation in the context of TNF α and

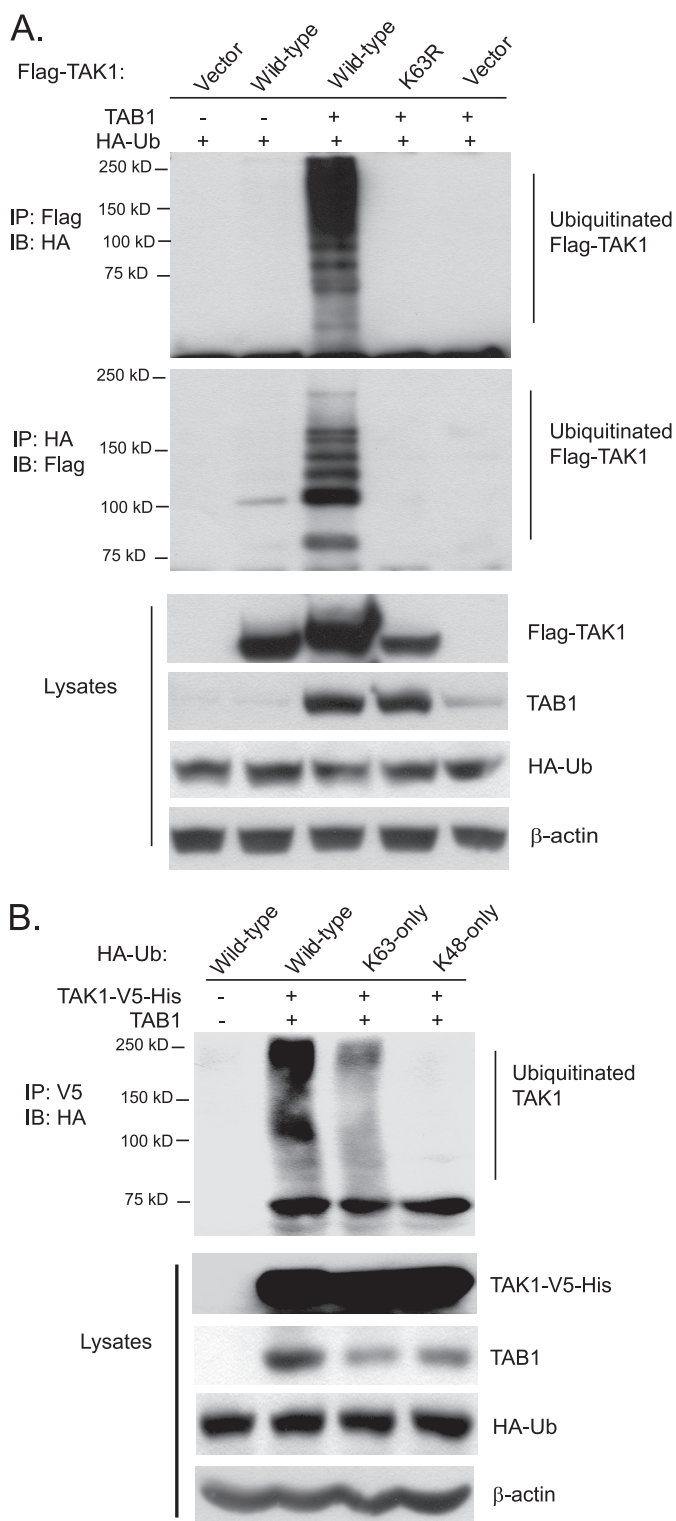


FIGURE 2. Co-overexpression of TAK1/TAB1 induces Lys⁶³-linked TAK1 polyubiquitination. *A*, co-overexpression of TAK1/TAB1 induces TAK1 polyubiquitination. Expression vectors encoding FLAG-TAK1 and HA-ubiquitin were co-transfected into HEK-293T cells with control vector and expression vectors encoding TAB1, respectively. FLAG-TAK1 proteins in the transfected cells were immunoprecipitated (IP) with anti-FLAG antibodies (or anti-HA antibodies) and immunoblotted (IB) with anti-HA antibodies (or anti-FLAG antibodies) to detect the presence of ubiquitinated FLAG-TAK1. *B*, co-overexpression of TAK1/TAB1 induces Lys⁶³-linked TAK1 polyubiquitination. Expression vectors encoding TAK1-V5-His and TAB1 were co-transfected into HEK-293T cells with control vector and expression vectors encoding HA-ubiquitin wild type, Lys⁶³-only, and Lys⁴⁸-only, respectively. TAK1-V5-His proteins

IL-1 β stimulation. Using biochemical and reporter assays, we identified the Lys¹⁵⁸ residue within TAK1 kinase domain as a Lys⁶³-linked polyubiquitination site required for TAK1-mediated IKK/NF- κ B and JNK/AP-1 activation. Furthermore, we found that polyubiquitination of Lys¹⁵⁸ residue can be induced by TNF α and IL-1 β stimulation and is required for TNF α - and IL-1 β -induced IKK γ /NEMO association with TAK1. In addition, we found that Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue is essential for TNF α and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation as well as IL-6 gene expression. Together, our results provide biochemical and genetic evidence that the TNF α - and IL-1 β -induced Lys⁶³-linked TAK1 polyubiquitination at Lys¹⁵⁸ is essential for TNF α and IL-1 β -induced TAK1 activation and TAK1-mediated IKK/NF- κ B and JNK/AP-1 activation.

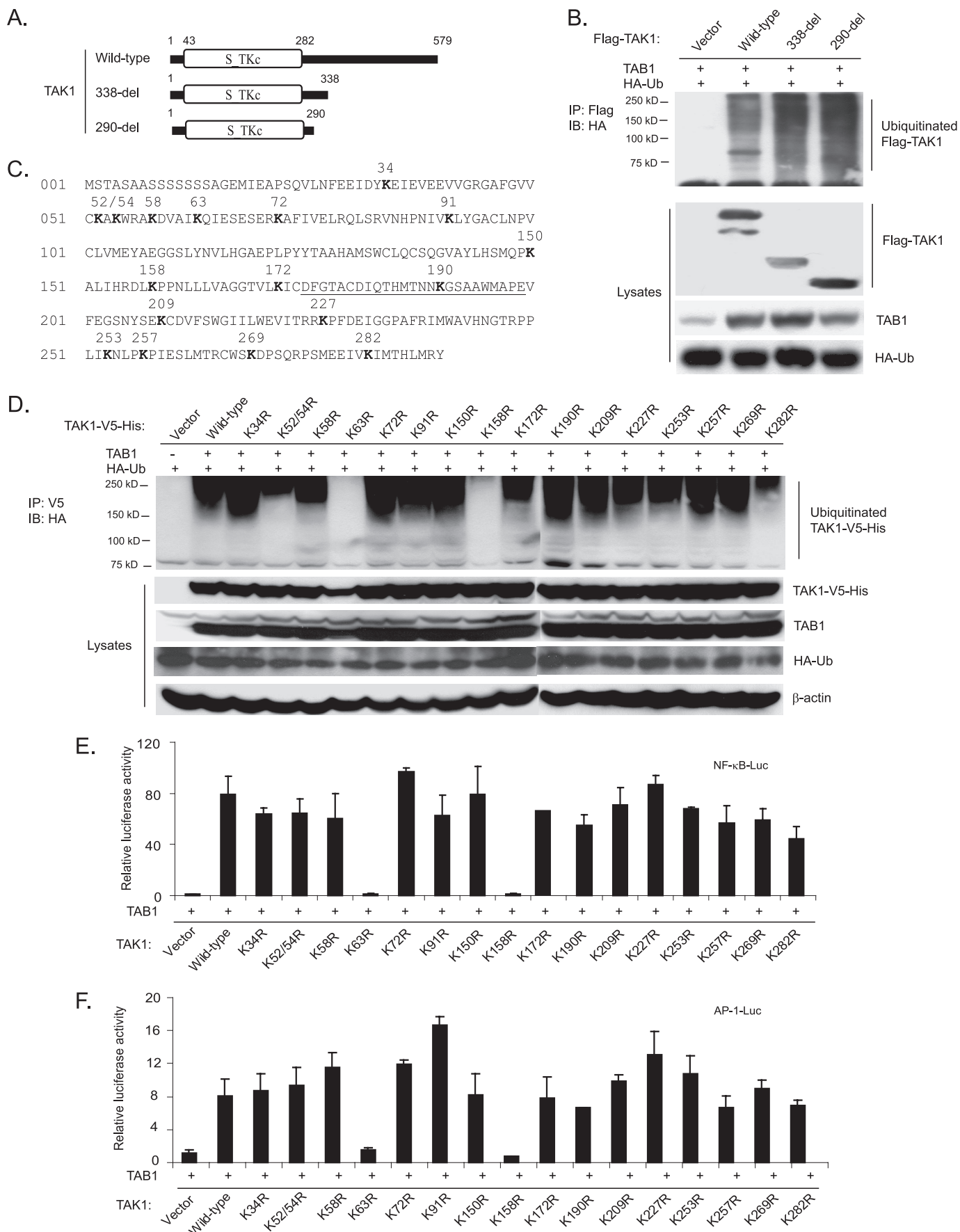
EXPERIMENTAL PROCEDURES

Purification of GST-tagged Fusion Proteins—Recombinant GST-TRAF2, GST-TRAF6, GST-TAK1-WT-V5-(1–292), GST-TAK1-K63R-V5-(1–292), GST-TAK1-K150R-V5-(1–292), GST-TAK1-K158R-V5-(1–292), and GST-TAB1 proteins were generated and purified as described previously (45). Briefly, the plasmids encoding recombinant proteins were transformed into *Escherichia coli* BL-21 strain (Invitrogen), and then recombinant protein expression was induced by 0.1 mM isopropyl β -D-thiogalactoside for 4 h at 30 °C. Bacteria were lysed with extraction buffer (50 mM Tris-HCl, pH 8.5, 100 mM NaCl, 1 mM EDTA, 1 mM dithiothreitol, 50 mg/ml lysozyme, 10 μ g/ml aprotinin, 10 μ g/ml leupeptin, and 1 mM phenylmethylsulphonyl fluoride) for 45 min on ice and sonicated at 4 °C in 1% Sarkosyl (Sigma). After adding Triton X-100 (1%), 5 μ g/ml DNase, and 5 μ g/ml RNase (Roche Applied Science), the lysates were centrifuged at 15,000 \times *g*, and the supernatants containing GST fusion protein were collected. Fusion proteins were purified from cell lysates using glutathione-Sepharose beads (Sigma) overnight at 4 °C. The beads were washed three times in extraction buffer containing 0.5% Triton X-100 and one time in extraction buffer containing 0.1% Triton X-100. Proteins were eluted in elution buffer (30.7% glutathione, 50 mM Tris-HCl, pH 8.0, 20% glycerol, 5 M NaCl) and dialyzed in 1 \times phosphate-buffered saline. The protein concentrations were then assessed with a Bradford protein assay (Bio-Rad). The proteins were resolved by 10% SDS-PAGE and visualized by Coomassie Blue staining of the gel.

In Vitro Kinase Assays—HEK-293T cells seeded onto 10-cm dishes were transfected with the TAK1-V5 and TAB1 expression plasmids. The TAK1-V5 proteins were immunoprecipitated from cell extracts with anti-V5 antibody and washed three times with the lysis buffer and twice with kinase reaction buffer (25 mM Tris, pH 7.5, 2 mM dithiothreitol, 0.1 mM Na₃VO₄, 10 mM MgCl₂). Then the immunoprecipitates were resuspended in kinase buffer containing 100 mM ATP. His-MKK6 fusion protein was added as a substrate for TAK1. After a 30-min incubation at 30 °C, the samples were separated by SDS-PAGE,

in the transfected cells were immunoprecipitated with anti-V5 antibodies and immunoblotted with anti-HA antibodies to detect the presence of ubiquitinated TAK1-V5-His.

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transferred to polyvinylidene difluoride membrane, and detected by anti-phospho-MKK6 antibodies.

In Vitro Ubiquitination Assays—For *in vitro* ubiquitination assays, recombinant GST proteins, including GST-TAK1-WT-V5-(1–292), GST-TAK1-K63R-V5-(1–292), GST-TAK1-K150R-V5-(1–292), GST-TAK1-K158R-V5-(1–292), GST-TAB1, GST-TRAF2, and GST-TRAF6, were added as indicated along with 50 nM UBE1, 875 nM E2 (Ubc13/Uev1a), 59 μ M ubiquitin (wild type, Lys⁶³-only, and Lys⁴⁸-only) and Mg-ATP (all from Boston Biochemicals) into a 20- μ l total volume of ubiquitination buffer (20 mM HEPES, pH 7.4, 1 mM MgCl₂, 1 mM dithiothreitol, 10 μ g/ml aprotinin, 10 μ g/ml leupeptin, 1 mM benzamide, 20 mM disodium *p*-nitrophenyl phosphate, 1 mM phenylmethylsulfonyl fluoride). The mixture was incubated at 30 °C for 2 h with gentle agitation. After stopping the reaction, the mixture was added with SDS to 1% concentration and then boiled for 10 min. Subsequently, the sample was diluted 10-fold with 1 \times phosphate-buffered saline before immunoprecipitation with anti-V5 antibodies and immunoblotting with anti-ubiquitin (Ub) antibodies.

RESULTS

TNF α and IL-1 β Induce Lys⁶³-linked TAK1 Polyubiquitination—TAK1 plays an essential role in the TNF α and IL-1 β -mediated IKK β and JNK activation (15, 31–33). Recent studies suggest that TAK1 activity is regulated by ubiquitination (15, 40, 42). However, the molecular details of ubiquitination in TAK1 activation remain poorly understood. To further understand the role of ubiquitination in TAK1 activation and examine whether TNF α and IL-1 β induce TAK1 ubiquitination, we generated a HeLa cell line with stable expression of human wild type TAK1 with an N-terminal FLAG tag in order to better detect the subtle change of TAK1 during TNF α - and IL-1 β -induced TAK1 activation. Then we stimulated FLAG-TAK1 HeLa cells with TNF α and IL-1 β for the time points indicated and lysed cells. FLAG-TAK1 proteins in the cell lysates were immunoprecipitated with anti-FLAG antibodies and immunoblotted with anti-ubiquitin antibodies. As shown in Fig. 1, A and B, TNF α and IL-1 β rapidly induced the ubiquitination of TAK1 and the induced TAK1 ubiquitination was diminished at the later time points after stimulation. To determine whether TNF α and IL-1 β induces Lys⁶³- or Lys⁴⁸-linked TAK1 polyubiquitination, we co-transfected FLAG-TAK1 into HEK-293T cells with HA-ubiquitin Lys⁶³ or Lys⁴⁸ only mutant and then stimulated cells with TNF α and IL-1 β for the times indicated. FLAG-TAK1 proteins in the cells were immunoprecipitated

with anti-FLAG antibodies and immunoblotted with anti-HA antibodies. In this assay, we found that TNF α and IL-1 β induced a much stronger Lys⁶³-linked TAK1 polyubiquitination within 5 min of stimulation compared with the Lys⁴⁸-linked TAK1 polyubiquitination (Fig. 1, C and D). These results demonstrate that TNF α and IL-1 β induce Lys⁶³-linked TAK1 polyubiquitination.

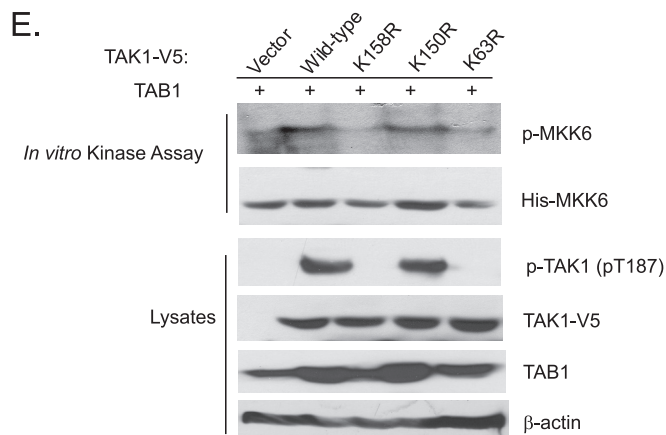
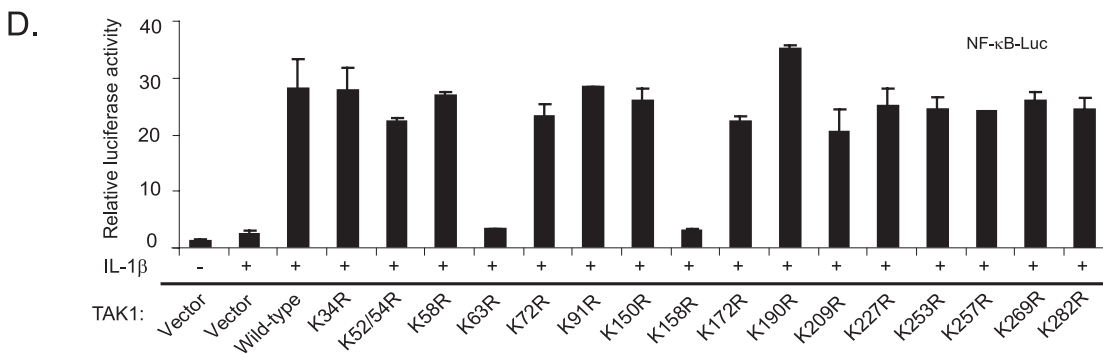
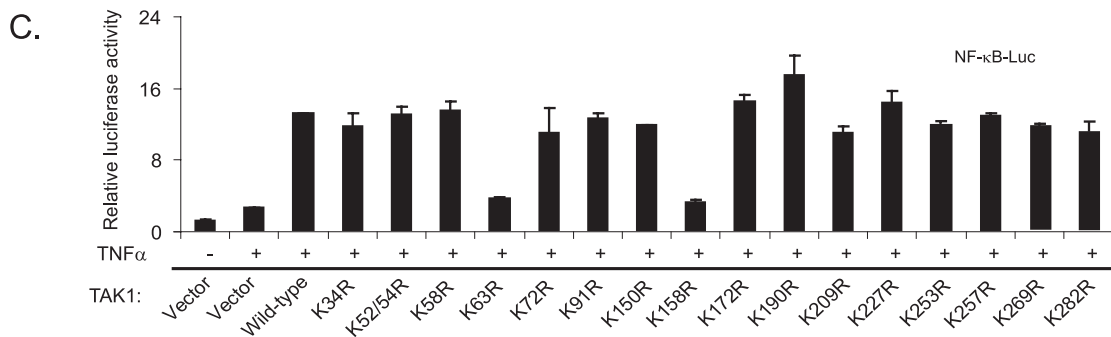
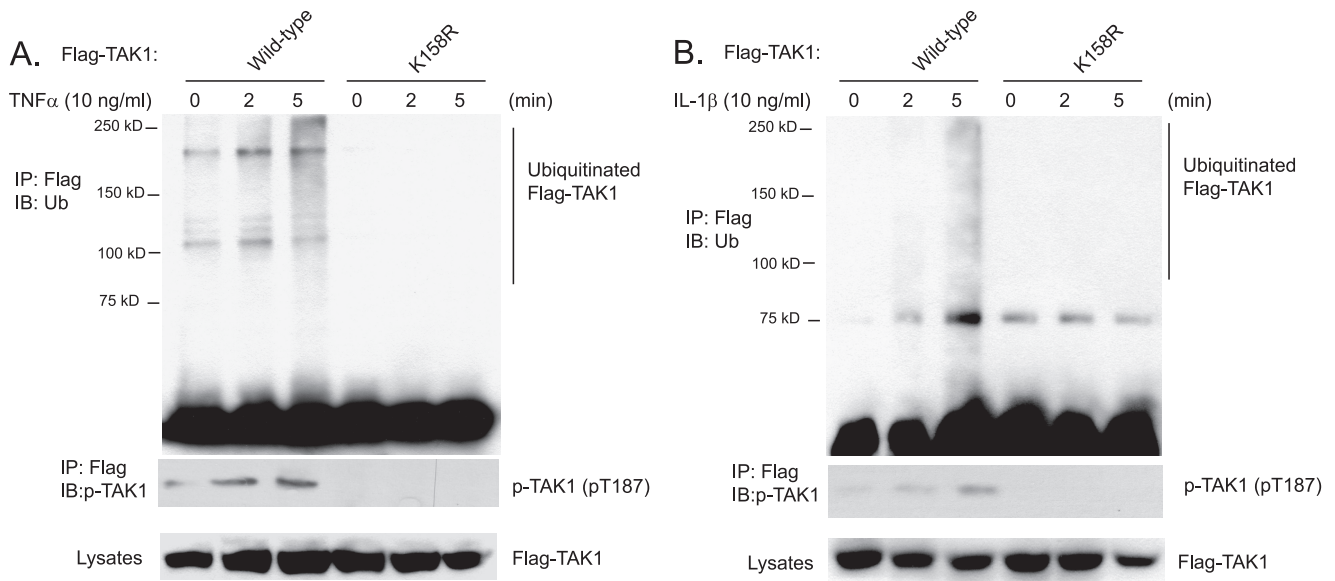
Co-overexpression of TAK1/TAB1 Induces Lys⁶³-linked TAK1 Polyubiquitination—Co-overexpression of TAK1 with its regulatory subunit TAB1 leads to TAK1 autophosphorylation and activation in the cells (36). Therefore, we hypothesized that co-overexpression of TAK1 with its regulatory subunit TAB1 might also lead to TAK1 ubiquitination. To test this hypothesis, we co-transfected HA-Ub expression vectors with or without FLAG-TAK1 and TAB1 expression vectors into HEK-293T cells. The cell lysates from the transfected cells were heated in the presence of 1% SDS and diluted with lysis buffer in order to disrupt noncovalent protein-protein interactions. Then FLAG-TAK1 or HA-ubiquitin in the cell lysates were immunoprecipitated with anti-FLAG or anti-HA antibodies and immunoblotted with anti-HA or anti-FLAG antibodies for the detection of the ubiquitinated TAK1. In this assay, we found that co-overexpression of TAK1 wild type with TAB1 induced TAK1 polyubiquitination, whereas co-overexpression of a kinase-dead TAK1-K63R mutant (TAK1 kinase ATP binding site K63R mutation) with TAB1 and overexpression of TAK1 alone both failed to do so (Fig. 2A). This result indicates that co-overexpression of TAK1 with TAB1 induces TAK1 polyubiquitination and that TAK1 polyubiquitination is associated with its activation.

To determine the linkage type of polyubiquitin chains on TAK1, we co-transfected TAK1-V5-His and TAB1 into HEK-293T cells along with HA-Ub wild type or mutant containing one lysine only at position 63 (Lys⁶³-only) or 48 (Lys⁴⁸-only). As shown in Fig. 2B, TAK1 polyubiquitination was observed with the Ub wild type and Lys⁶³-only mutant but not with the Lys⁴⁸-only mutant. This result suggests that polyubiquitin chains on TAK1 were linked primarily through Lys⁶³ of Ub. Together, these results indicate that co-overexpression of TAK1 with TAB1 induces Lys⁶³-linked TAK1 polyubiquitination and that Lys⁶³-linked TAK1 polyubiquitination is associated with its activation.

Co-overexpression of TAK1/TAB1 Induces Lys⁶³-linked TAK1 Polyubiquitination at the Lys¹⁵⁸ Residue within the Kinase Domain—To address the role of the TAK1 Lys⁶³-linked polyubiquitination in its activation and TAK1-mediated

FIGURE 3. Co-overexpression of TAK1/TAB1 induces Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue within the kinase domain. A, schematic representation of TAK1 wild type and deletion mutants with the kinase domain indicated. B, co-overexpression of the TAK1/TAB1-induced TAK1 polyubiquitination site is located within the kinase domain. Expression vectors encoding HA-ubiquitin and TAB1 were co-transfected into HEK-293T cells with control vector and expression vectors encoding FLAG-TAK1 wild type and two deletion mutants (338-del and 290-del), respectively. FLAG-TAK1 proteins in the transfected cells were immunoprecipitated (IP) with anti-FLAG antibodies and immunoblotted (IB) with anti-HA antibodies to detect the presence of ubiquitinated FLAG-TAK1. C, TAK1 primary sequence with the lysine residues within its N-terminal 290 amino acids indicated. The kinase activation loop is underlined. D, co-overexpression of TAK1/TAB1 induces TAK1 polyubiquitination at Lys¹⁵⁸ within the kinase domain. Expression vectors encoding HA-ubiquitin and TAB1 were co-transfected into HEK-293T cells with control vector and expression vectors encoding TAK1-V5-His wild type and 16 lysine to arginine mutants, respectively. TAK1-V5-His proteins in the transfected cells were immunoprecipitated with anti-V5 antibodies and immunoblotted with anti-HA antibodies to detect the presence of ubiquitinated TAK1-V5-His. E and F, the effect of overexpression of TAK1 lysine to arginine mutants with TAB1 on TAK1/TAB1-induced NF- κ B (E) and AP-1 (F) activation. TAB1 expression vectors, NF- κ B luciferase reporter, and control *Renilla* luciferase reporter vectors were co-transfected into TAK1-deficient MEF cells with empty vector or expression vectors encoding TAK1 wild type and lysine to arginine mutants, respectively. The relative luciferase activity was measured 48 h later and normalized with the *Renilla* activity. Error bars, \pm S.D. in triplicate experiments.

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downstream signaling events, it is essential to map the Ub lysine acceptor site(s) of TAK1 and characterize the functional effects of eliminating the site(s). To determine the location of the Ub lysine acceptor site(s) of TAK1, we first generated two TAK1 C-terminal deletion mutants that sequentially deleted the TAK1 regulatory domain at amino acid 338 and 290, respectively (Fig. 3A). Then we co-transfected HA-Ub and TAB1 into HEK-293T cells along with TAK1 wild type or two TAK1 C-terminal deletion mutants (338-del and 290-del). As shown in Fig. 3B, TAK1 wild type and two TAK1 C-terminal deletion mutants were polyubiquitinated at similar levels. Therefore, we hypothesized that one or more Lys residues within the TAK1 N-terminal kinase domain (1–290 amino acids) may serve as the Ub lysine acceptor site(s).

TAK1 contains 17 lysine residues within the first 290 amino acids, including its kinase domain (Fig. 3C). To identify the likely Ub lysine acceptor site(s) in TAK1, we systematically replaced each of these Lys residues with an Arg residue, which maintains the positive charge but does not serve as an acceptor site for Ub modification. The expression vectors encoding the C-terminal V5-tagged TAK1 wild type and mutants with different Lys to Arg mutations were co-transfected with TAB1 into HEK-293T cells along with HA-Ub. The cell lysates were boiled in the presence of 1% SDS and immunoprecipitated with anti-V5, followed by SDS-PAGE and immunoblotting with anti-HA antibodies. Of the 17 Lys to Arg mutants examined, only K63R (a kinase ATP binding site mutant) and K158R mutants failed to induce TAK1 polyubiquitination compared with the other mutants (Fig. 3D).

Consistent with these results, luciferase analysis with NF- κ B and AP-1-responsive reporters showed that TAK1 K63R and K158R mutants failed to induce the luciferase reporter gene expression, whereas TAK1 wild type and other mutants induced a high level of reporter gene activity when co-overexpressed with TAB1 in TAK1-deficient MEF cells (Fig. 3, E and F). Because the TAK1 Lys⁶³ residue is a conserved lysine residue in the ATP pocket essential for coordinating kinase ATP binding and activation, it is likely that Lys¹⁵⁸ is the predominant Ub acceptor site for TAK1 polyubiquitination. Taken together, these results suggest that polyubiquitination of TAK1 at Lys¹⁵⁸ is required for the TAK1/TAB1 co-overexpression-induced NF- κ B and AP-1 activation.

The Lys¹⁵⁸ Residue Is Required for TNF α - and IL-1 β -induced TAK1 Polyubiquitination and NF- κ B Activation—To determine whether the Lys¹⁵⁸ residue is an acceptor site for Ub modification of TAK1 in response to TNF α and IL-1 β stimulation, we stimulated the stable HeLa cell lines harboring FLAG-TAK1 wild type and FLAG-TAK1 K158R mutant with TNF α and

IL-1 β . As shown in Fig. 4, A and B, FLAG-TAK1 polyubiquitination and phosphorylation were rapidly induced by TNF α and IL-1 β in the cells expressing FLAG-TAK1 wild type but not K158R mutant. This result suggests that the Lys¹⁵⁸ residue is an acceptor site for Ub modification of TAK1 in response to TNF α and IL-1 β stimulation.

To determine whether the TAK1 Lys¹⁵⁸ residue is required for TNF α - and IL-1 β -induced NF- κ B activation, we performed luciferase analysis in TAK1-deficient MEF cells transfected with TAK1 wild type and various lysine to arginine mutants with NF- κ B-responsive reporter. In these assays, TAK1 K63R and K158R mutants blocked the TNF α and IL-1 β -induced NF- κ B-dependent luciferase activities, whereas TAK1 wild type and other lysine to arginine mutants mediated a high level of reporter gene activity (Fig. 4, C and D). To further confirm the role of Lys⁶³-linked polyubiquitination at Lys¹⁵⁸ in TAK1 activation, we co-transfected TAB1 into HEK-293T cells with vector control, TAK1-V5 wild type, K158R, K150R, and K63R, respectively. TAK1 wild type and mutant proteins were immunoprecipitated with anti-V5 antibodies for an *in vitro* kinase assay using recombinant MKK6 as a substrate. In this assay, we found that TAK1 K158R and K63R mutants failed to phosphorylate MKK6 *in vitro* compared with TAK1 wild type and K150R mutant (Fig. 4E). Taken together, these results suggest that polyubiquitination of TAK1 at Lys¹⁵⁸ residue is required for TNF α - and IL-1 β -induced NF- κ B activation.

TRAF2 and TRAF6 Mediate Lys⁶³-linked TAK1 Polyubiquitination at the Lys¹⁵⁸ Residue *In Vivo* and *In Vitro*—TRAF2 and TRAF6 have been identified to act as the Ub E3 ligases in the TNF α - and IL-1 β -mediated NF- κ B signal transduction pathway to facilitate the synthesis of unique Lys⁶³-linked poly-Ub chains rather than the conventional Lys⁴⁸-linked poly-Ub chains that target proteins for degradation (46). To determine whether TRAF2 and TRAF6 mediate Lys⁶³-linked TAK1 polyubiquitination, we co-transfected TAK1-V5-His and TRAF2/6 into HEK-293T cells along with HA-Ub wild type, Lys⁴⁸-only, and Lys⁶³-only, respectively. As shown in Fig. 5, A and B, overexpression of TRAF2 and TRAF6 induced a strong TAK1 polyubiquitination with Ub wild type and Lys⁶³-only but not Lys⁴⁸-only chains. This result suggests that TRAF2 and TRAF6 are the Ub E3 ligases that mediate Lys⁶³-linked polyubiquitination of TAK1 in the cells.

To further determine whether TRAF2 and TRAF6 are the Ub E3 ligases that mediate Lys⁶³-linked polyubiquitination of TAK1, we set up an *in vitro* ubiquitination assay with purified ubiquitin wild type, Lys⁴⁸-only and Lys⁶³-only mutants as well as GST-TRAF2, GST-TRAF6, Ubc13/Uev1A, E1/ATP, GST-TAB1, GST-TAK1-(1–292) wild type, and K150R and K158R

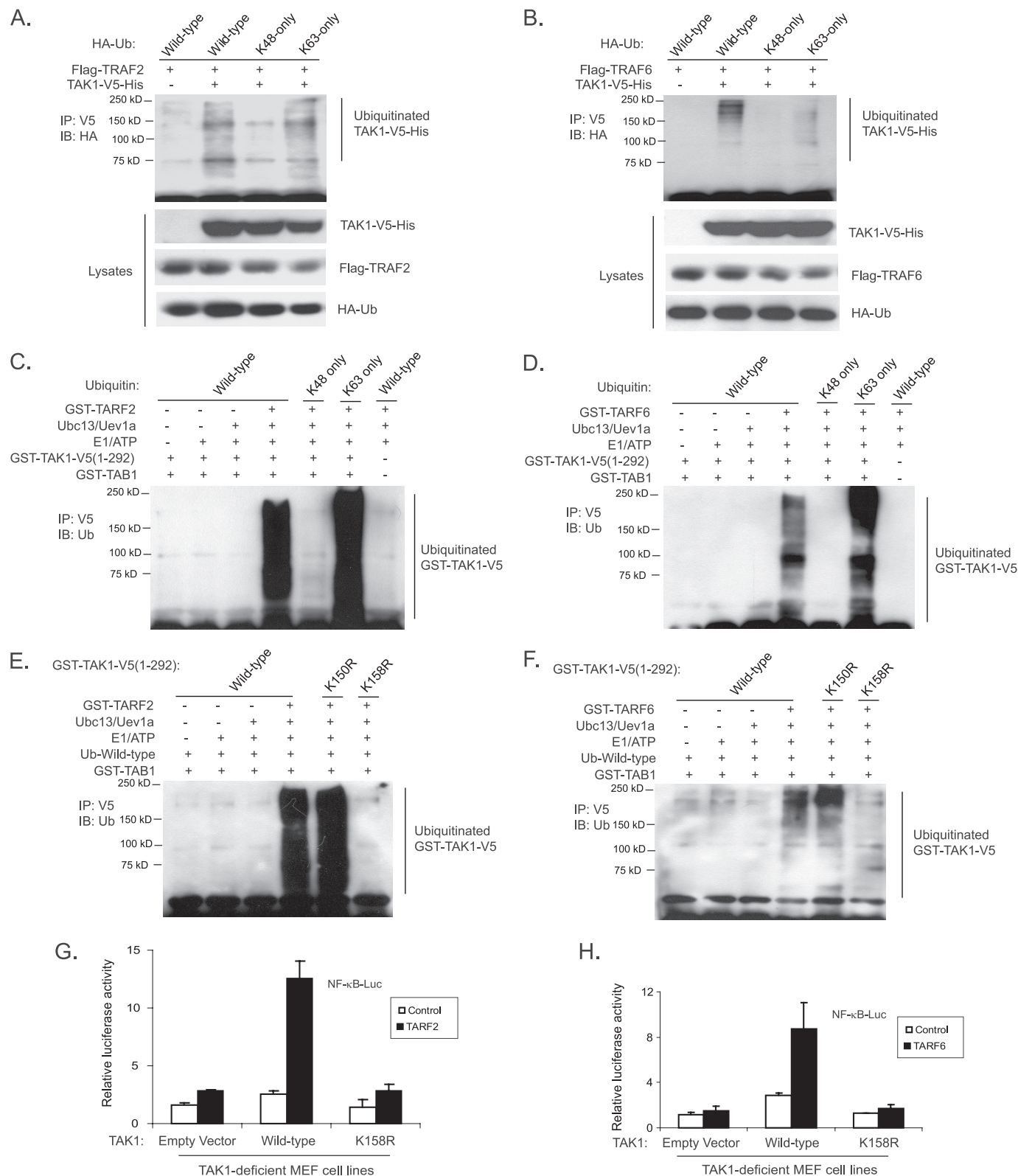
FIGURE 4. TNF α and IL-1 β induce TAK1 polyubiquitination at the Lys¹⁵⁸ residue. A and B, TNF α and IL-1 β induce TAK1 polyubiquitination at Lys¹⁵⁸. HeLa cells with stable expression of FLAG-TAK1 wild type and K158R mutant were either untreated or treated with TNF α (10 ng/ml) (A) and IL-1 β (10 ng/ml) (B) for the time points indicated and subsequently lysed. FLAG-TAK1 proteins in the cell lysates were immunoprecipitated (IP) with anti-FLAG antibodies and immunoblotted (IB) with anti-ubiquitin antibodies to detect the presence of ubiquitinated FLAG-TAK1. C and D, the effect of overexpression of TAK1 lysine to arginine mutants on TNF α -induced (C) and IL-1 β -induced (D) NF- κ B activation. NF- κ B luciferase reporter and control *Renilla* luciferase reporter vectors were co-transfected into TAK1-deficient MEF cells with empty vector or expression vectors encoding TAK1 wild type and lysine to arginine mutants for 48 h, respectively. Cells were then either untreated or treated with TNF α (1 ng/ml) and IL-1 β (1 ng/ml) for 6 h. The relative luciferase activity was measured and normalized with the *Renilla* activity. The error bars indicate \pm S.D. in triplicate experiments. E, TAK1 Lys¹⁵⁸ is required for TAK1 activation. TAB1 were transfected into HEK-293T cells with TAK1-V5 wild type, K158R, K150R, and K63R, respectively. TAK1-V5 proteins were immunoprecipitated from cell extracts with anti-V5 antibodies for an *in vitro* kinase assay using recombinant His-MKK6 as a substrate.

TNF and IL-1 Induce TAK1 Polyubiquitination at Lys¹⁵⁸

mutants, respectively. As shown in Fig. 5, C and D, both TRAF2 and TRAF6 catalyze the Ub wild type and Lys⁶³-linked but not Lys⁴⁸-linked TAK1 polyubiquitination. Notably, recombinant TRAF2 and TRAF6 mediated a strong polyubiquitination of TAK1 wild type and K150R mutant, whereas they failed to

mediate polyubiquitination of TAK1 K158R mutant completely (Fig. 5, E and F).

To further confirm that TAK1 K158R is the predominant Ub acceptor site, we performed *in vitro* ubiquitination assays with GST-TAK1-(1–292) wild type and K63R and K158R mutants,



respectively. In these assays, we found that TRAF2 and TRAF6 catalyze a strong polyubiquitination of TAK1 wild type and K63R mutant, whereas they failed to mediate polyubiquitination of the TAK1 K158R mutant (supplemental Fig. S1). These results suggest that TRAF2 and TRAF6 act as the Ub E3 ligases to mediate Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue.

Consistent with these results, luciferase analysis with NF- κ B- and AP-1-responsive reporters showed that TAK1 K158R mutant but not wild type blocked TRAF2 or TRAF6 overexpression-induced luciferase reporter gene activity (Fig. 5, G and H). This result indicates that Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue is required for TRAF2- and TRAF6-mediated NF- κ B activation.

UBC13 has been shown to play an important role in TRAF6-mediated Lys⁶³-linked polyubiquitination (8, 15). To further confirm the role of the Lys⁶³-linked TAK1 polyubiquitination in TAK1-mediated NF- κ B activation, we cotransfected NF- κ B reporter and TAK1/TAB1 into HEK-293T cells along with sh-Control or shUBC13 plasmids. In this reporter assay, we found that suppression of UBC13 expression inhibited TAK1/TAB1-induced NF- κ B activation (supplemental Fig. S2). Taken together, these results demonstrate that TRAF2 and TRAF6 are the Ub E3 ligases that specifically catalyze Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue in the cells.

IKK γ /NEMO Binds to Polyubiquitinated TAK1—The regulatory subunit IKK γ /NEMO of the IKK complex binds polyubiquitin chains and mediates TNF α - and IL-1 β -induced IKK β -NF- κ B activation (7, 25). Previous studies demonstrate that TAK1 is the major kinase mediating TNF α and IL-1 β -induced IKK/NF- κ B activation (15, 31–33). Therefore, we hypothesized that IKK γ /NEMO binds to Lys⁶³-linked polyubiquitinated TAK1 for IKK activation. To test this hypothesis, we co-transfected TAK1 wild type or K158R mutant with HA-Ub and TAB1 into HEK-293T cells and subsequently lysed the cells. Then we incubated recombinant GST and GST-IKK γ /NEMO bound to glutathione-agarose beads with the above cell lysates for the detection of the amount of TAK1 that was pulled down. As shown in Fig. 6A, GST-IKK γ /NEMO but not GST control pulled down a significant amount of polyubiquitinated TAK1 wild type but not polyubiquitination-deficient K158R mutant proteins. This result suggests that IKK γ /NEMO binds to the polyubiquitinated TAK1.

To determine whether IKK γ /NEMO binds to polyubiquitinated TAK1 in the cells in response to TNF α and IL-1 β stimu-

lation, we stimulated HeLa cell lines expressing FLAG-TAK1 wild type and K158R mutant with TNF α and IL-1 β for the times indicated, and endogenous NEMO was immunoprecipitated from the cell lysates and immunoblotted with anti-FLAG antibodies to detect the presence of FLAG-TAK1. As shown in Fig. 6, B and C, the ubiquitinated FLAG-TAK1 wild type but not ubiquitination-deficient K158R was rapidly pulled down by immunoprecipitation of IKK γ /NEMO in response to TNF α and IL-1 β stimulation.

To further determine the role of TAK1 Lys¹⁵⁸ in TNF α - and IL-1 β -induced complex formation, including TRAFs, TAK1/TAB1, and IKKs, we transfected expression vectors encoding wild type and K158R TAK1-V5-His into HEK-293T cells and stimulated cells with TNF α and IL-1 β (IL-1 β receptor expression vectors were co-transfected in the IL-1 β -stimulated cells) at the time points indicated. TAK1-V5-His proteins in the cell lysates were pulled down with nickel affinity beads and immunoblotted with anti-TRAF2/6 and anti-IKK β antibodies. In this assay, we found that TNF α and IL-1 β induced association of TRAF2 and -6 with both TAK1 wild type and K158R mutant equally well, whereas TNF α and IL-1 β failed to induce significant interaction of TAK1 K158R mutant with IKK β compared with TAK1 wild type (supplemental Fig. S3). Taken together, these results demonstrate that TNF α - and IL-1 β -induced polyubiquitination of TAK1 is required for the association of TAK1 with IKK γ /NEMO and IKK β .

TAK1 Polyubiquitination at the Lys¹⁵⁸ Residue Is Required for TNF α - and IL-1 β -induced IKK/NF- κ B and JNK/AP-1 Activation—To further explore the role of polyubiquitination of TAK1 at the Lys¹⁵⁸ residue in TNF α and IL-1 β -induced IKK/NF- κ B and JNK/AP-1 activation, the TAK1 wild type and K158R mutant expression vectors were stably introduced back into the TAK1-deficient MEF cells by a retroviral transduction system. Then TAK1-deficient, TAK1 wild type, and K158R mutant reconstituted MEF cells were treated with TNF α or IL-1 β at different time points, and then the cell lysates were immunoblotted with the indicated antibodies to examine TNF α - and IL-1 β -induced JNK, p38 MAPK, IKK, and I κ B α phosphorylation as well as I κ B α degradation. In this assay, TNF α - and IL-1 β -induced-JNK2 and p38 phosphorylation were completely blocked in the TAK1-deficient and K158R mutant MEF cells, whereas TNF α and IL-1 β -induced-JNK1 activation was only partially inhibited in the TAK1-deficient and K158R mutant MEF cells compared with the TAK1 wild type reconstituted cells (Fig. 7, A and B). Similarly, TNF α and

FIGURE 5. TRAF2 and TRAF6 mediate Lys⁶³-linked TAK1 polyubiquitination at the Lys¹⁵⁸ residue *in vivo* and *in vitro*. A and B, overexpression of TRAF2 (A) and TRAF6 (B) induces Lys⁶³-linked TAK1 polyubiquitination. Expression vectors encoding TAK1-V5-His and TRAF2 (A) or TRAF6 (B) were co-transfected into HEK-293T cells with control vector and expression vectors encoding HA-ubiquitin wild type, Lys⁶³-only and Lys⁴⁸-only, respectively. TAK1-V5-His proteins in the transfected cells were immunoprecipitated (IP) with anti-V5 antibodies and immunoblotted with anti-ubiquitin antibodies to detect the presence of ubiquitinated TAK1-V5-His. C and D, TRAF2 (C) and TRAF6 (D) mediate Lys⁶³-linked TAK1 polyubiquitination *in vitro*. Recombinant GST-TAK1-V5-(1–292), GST-TAB1, E1, E2, (Ubc13/Uev1a), and GST-TRAF2 (C) or GST-TRAF6 (D) were co-incubated in the ubiquitination buffer with recombinant ubiquitin wild type, Lys⁴⁸-only, and Lys⁶³-only, respectively. After 2 h, GST-TAK1-V5-(1–292) proteins were immunoprecipitated with anti-V5 antibodies and immunoblotted with anti-ubiquitin antibodies to detect the presence of ubiquitinated TAK1-V5-(1–292). E and F, TRAF2 (E) and TRAF6 (F) mediate Lys⁶³-linked TAK1 polyubiquitination at Lys¹⁵⁸ *in vitro*. Recombinant ubiquitin wild type, GST-TAB1, E1, E2 (Ubc13/Uev1a), and GST-TRAF2 (E) or GST-TRAF6 (F) were co-incubated in the ubiquitination buffer with GST-TAK1-V5-(1–292) wild type, K158R, or K150R, respectively. After 2 h, GST-TAK1-V5-(1–292) wild type and mutant proteins were immunoprecipitated with anti-V5 antibodies and immunoblotted with anti-ubiquitin antibodies to detect the presence of ubiquitinated TAK1-V5-(1–292). G and H, TAK1 lysine 158 to arginine mutant inhibits TRAF2-induced (G) and TRAF6-induced (H) NF- κ B activation. TRAF2 or TRAF6 expression vectors, NF- κ B luciferase reporter, and control *Renilla* luciferase reporter vectors were co-transfected into TAK1-deficient MEF cells with empty vector or expression vectors encoding TAK1 wild type and lysine 158 to arginine (K158R) mutant. The relative luciferase activity was measured 48 h later and normalized with the *Renilla* activity. Error bars, \pm S.D. in triplicate experiments.

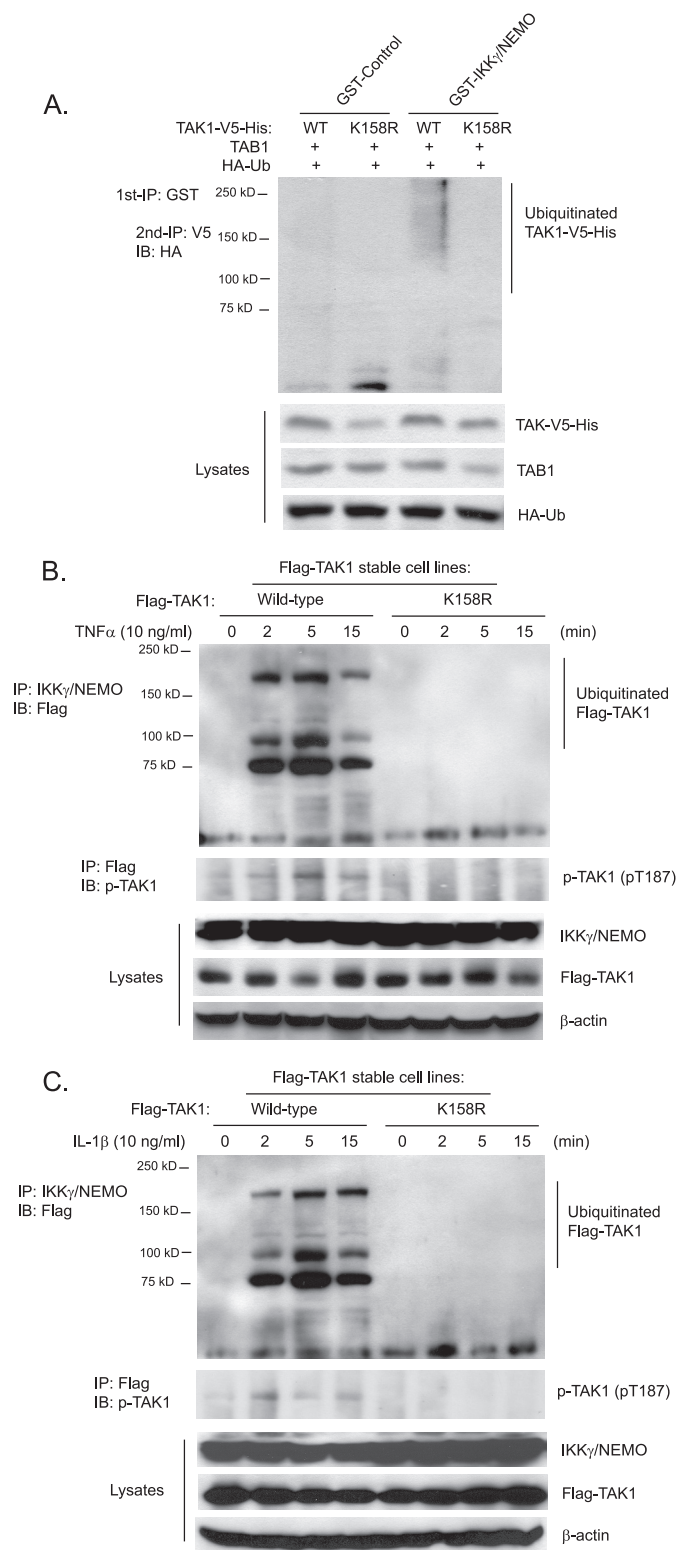


FIGURE 6. IKK γ /NEMO binds to polyubiquitinated TAK1. *A*, recombinant IKK γ /NEMO pulls down polyubiquitinated TAK1. Expression vectors encoding TAB1 and HA-ubiquitin were co-transfected into HEK-293T cells with expression vectors encoding TAK1-V5-His wild type and K158R mutant, respectively. The cell lysates were co-incubated with recombinant GST control and GST-IKK γ /NEMO proteins bound to glutathione-agarose beads for 2 h, respectively. GST control and GST-IKK γ /NEMO proteins bound to glutathione-agarose beads were first precipitated with centrifugation and boiled. Then co-precipitated TAK1-V5-His proteins were immunoprecipitated (IP) with anti-V5 antibodies and immunoblotted (IB) with anti-HA antibodies to detect the presence of polyubiquitinated TAK1-V5-His proteins. *B* and *C*,

IL-1 β -induced IKK and I κ B α phosphorylation as well as I κ B α degradation were impaired in the MEF cells with control vector and TAK1 K158R mutant compared with the cells with TAK1 wild type (Fig. 7, *A* and *B*). We also isolated nuclear extracts from the above cells treated with TNF α and IL-1 β at the indicated time points and found that TNF α - and IL-1 β -induced NF- κ B nuclear translocation was also significantly impaired in the MEF cells with control vector and TAK1 K158R mutant compared with cells with TAK1 wild type (Fig. 7, *A* and *B*). These results suggest that the TAK1 Lys¹⁵⁸ residue is required for TNF α and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation.

Consistent with the above results, TNF α and IL-1 β induced much higher NF- κ B- and AP-1-dependent luciferase activities in MEF cells expressing the TAK1 wild type, compared with the TAK1-deficient and K158R mutant cells (Fig. 7, *C–F*).

Consistent with earlier reports that TAK1 is required for lipopolysaccharide-induced but not lysophosphatidic acid and protein kinase C-induced IKK/NF- κ B activation (15, 32, 33, 47), we found that TAK1 K158R mutant inhibited lipopolysaccharide-mediated but not lysophosphatidic acid- and protein kinase C-mediated IKK/NF- κ B activation (supplemental Figs. S4 and S5).

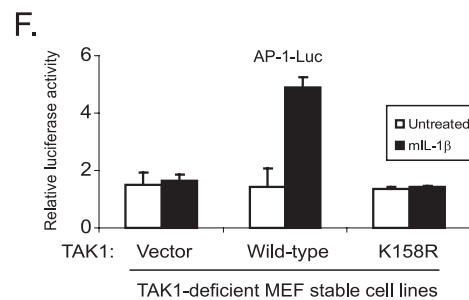
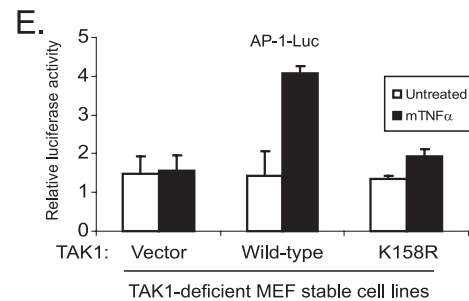
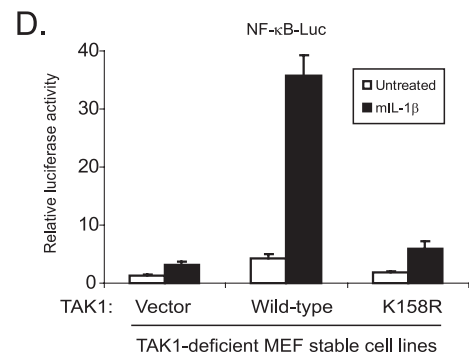
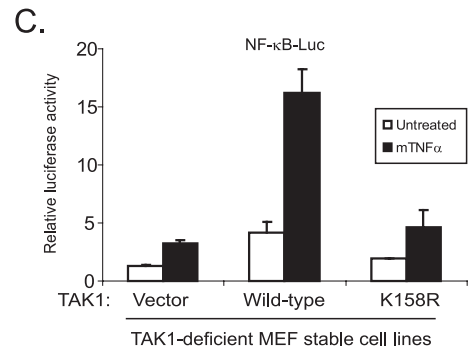
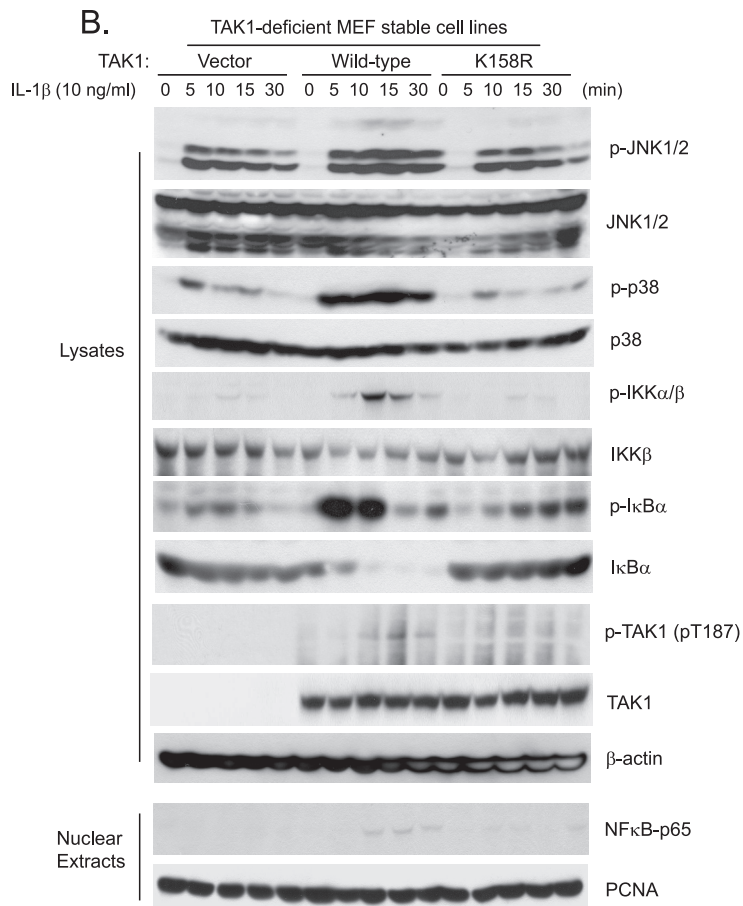
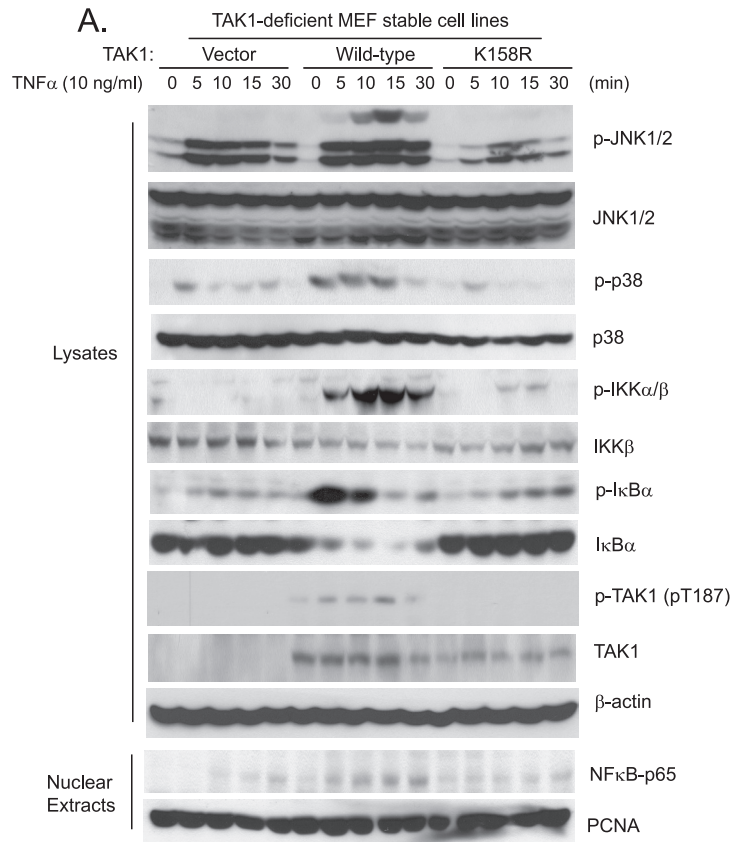
Taken together, these results indicate that polyubiquitination of TAK1 at the Lys¹⁵⁸ site is required for the TNF α - and IL-1 β -induced optimal IKK/NF- κ B and JNK/AP-1 activation.

TAK1 Polyubiquitination at the Lys¹⁵⁸ Residue Is Required for TNF α - and IL-1 β -induced IL-6 Gene Expression—TAK1 mediates TNF α - and IL-1 β -induced NF- κ B and AP-1 activation and IL-6 gene expression (48, 49). To determine the role of TAK1 polyubiquitination at the Lys¹⁵⁸ residue in TNF α - and IL-1 β -induced IL-6 gene expression, the TAK1-deficient stable MEF cell lines with vector control, TAK1 wild type, and K158R mutant were treated with TNF α or IL-1 β . Then we examined the TNF α - and IL-1 β -induced IL-6 gene expression at the mRNA level in the cells. As shown in Fig. 8, *A* and *B*, TNF α - and IL-1 β -induced IL-6 expression was blocked in the TAK1-deficient and K158R mutant cells compared with the TAK1 wild type cells. Consistent with this result, TNF α - and IL-1 β -induced IL-6 production in the cell medium was significantly inhibited in the TAK1-deficient and K158R mutant cells compared with the TAK1 wild type cells (Fig. 8, *C* and *D*). These results demonstrate that polyubiquitination at the Lys¹⁵⁸ residue is required for TNF α - and IL-1 β -induced IL-6 expression and production in the cells.

DISCUSSION

In this report, we further examined the mechanism of ubiquitination in TNF α - and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation and gained novel insights into the regulatory role of Lys⁶³-linked polyubiquitination in the TAK1-mediated

TNF α (*B*) and IL-1 β (*C*) induce association of IKK γ /NEMO with TAK1 wild type but not K158R mutant proteins. HeLa cells with stable expression of FLAG-TAK1 wild type and K158R mutant were either untreated or treated with TNF α (10 ng/ml) (*B*) and IL-1 β (10 ng/ml) (*C*) for the times indicated and subsequently lysed. Endogenous IKK γ /NEMO proteins in the cell lysates were immunoprecipitated with anti-NEMO antibodies and immunoblotted with anti-FLAG antibodies to detect the co-immunoprecipitation of FLAG-TAK1.



TNF and IL-1 Induce TAK1 Polyubiquitination at Lys¹⁵⁸

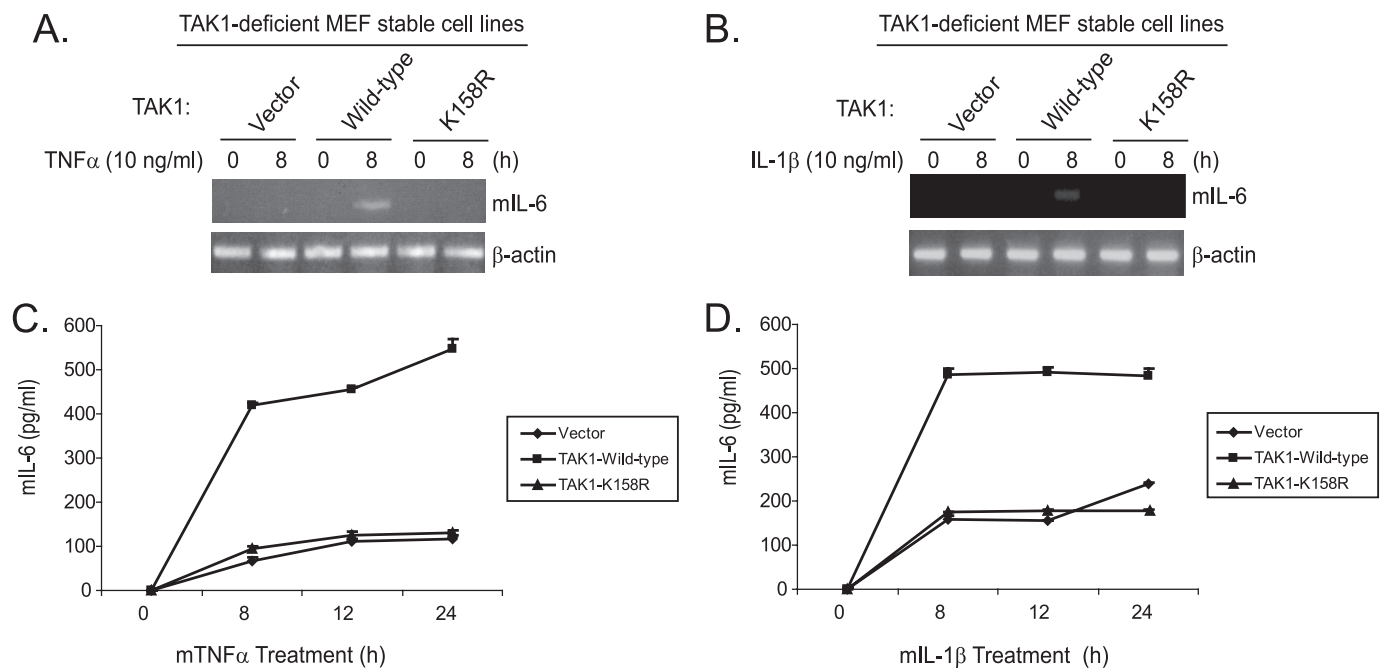


FIGURE 8. Polyubiquitination of TAK1 at Lys¹⁵⁸ residue is required for TNF α and IL-1 β -induced IL-6 gene expression. A and B, expression of the TAK1 K158R mutant inhibits IL-1-induced IL-6 gene transcription. TAK1-deficient, TAK1 wild type, and K158R mutant reconstituted MEF cells were untreated or treated with TNF α (10 ng/ml) (A) and IL-1 β (10 ng/ml) (B) for 8 h and subsequently harvested for extraction of total RNA using TRIzol reagent. 1 μ g of total RNA was used to synthesize first-strand cDNA using a reverse transcription kit according to the manufacturer's instructions. These synthesized cDNAs were used as templates for mouse IL-6 PCR amplification. The PCR products were resolved in 2% agarose gel. C and D, expression of the TAK1 K158R mutant inhibits IL-1-induced IL-6 production. TAK1-deficient, TAK1 wild type, and K158R mutant reconstituted MEF cells were untreated or treated with TNF α (2 ng/ml) (C) and IL-1 β (2 ng/ml) (D) for the times indicated. The supernatants from the cell cultures were collected and subjected to the mouse IL-6 ELISA analysis according to the manufacturer's instruction.

NF- κ B and AP-1 activation. Here we show that TAK1 is polyubiquitinated with a Lys⁶³-linked ubiquitin chain at the Lys¹⁵⁸ residue within its kinase domain in response to TNF α and IL-1 β stimulation and that Lys⁶³-linked polyubiquitination of TAK1 at the Lys¹⁵⁸ residue binds to IKK γ /NEMO, resulting in the recruitment of IKK complexes to the activated TAK1. Furthermore, we show that TRAF2 and TRAF6 are the Ub E3 ligases that mediate Lys⁶³-linked polyubiquitination of TAK1. Interestingly, a lysine to arginine mutation at Lys¹⁵⁸ of TAK1 blocks TNF α - and IL-1 β -induced TAK1 polyubiquitination and downstream IKK/NF- κ B and JNK/AP-1 activation. Taken together, our results indicate that Lys⁶³-linked polyubiquitination of TAK1 at Lys¹⁵⁸ and binding of polyubiquitinated TAK1 by IKK γ /NEMO are required for the TNF α - and IL-1-mediated IKK activation. In addition to the polyubiquitin chains on RIP1, TRAF2/6 and IRAKs, the polyubiquitin chains on TAK1 also serve as a part of a platform to recruit the IKK complex, JNK, and p38 MAPKs, allowing TAK1 to activate IKK, JNK, and p38-mediated signal transductions. We propose that Lys⁶³-linked polyubiquitination of TAK1 at Lys¹⁵⁸ within the kinase domain

is a general mechanism for TAK1-mediated IKK/NF- κ B and JNK/AP-1 activation.

TRAF2/RIP1 and TRAF6/IRAKs are proximal signaling components in TNF α and IL-1 β receptor-mediated IKK/NF- κ B and JNK/AP-1 activation, respectively (6, 7). Both TRAF2 and TRAF6 contain a RING domain that confers their E3 activity (46). Our studies demonstrate that TRAF2 and TRAF6 specifically catalyze TAK1 ubiquitination at the Lys¹⁵⁸ residue *in vivo* and *in vitro*. Considering there are 34 lysine residues in TAK1 protein sequence, it is very unusual that TRAF2- and TRAF6-mediated TAK1 polyubiquitination is so site-specific and consistent in both *in vivo* and *in vitro* ubiquitination assays. In the case of RIP1, none of the lysine mutations prevented RIP1 ubiquitination or NF- κ B activation once over-expressed in HEK-293T cells (7).

Previous studies suggest that TAK1 as a serine/threonine kinase is recruited to receptor associated TRAF2/RIP1 and TRAF6/IRAKs through binding of TAB2 and TAB3 to polyubiquitinated RIP1 and IRAKs in response to TNF α and IL-1 β stimulation, whereas the IKK complex is recruited via binding

FIGURE 7. Polyubiquitination of TAK1 at Lys¹⁵⁸ residue is required for TNF α - and IL-1 β -induced optimal IKK/NF- κ B and JNK/AP-1 activation. A and B, expression of the TAK1 K158R mutant inhibits TNF α -induced (A) and IL-1 β -induced (B) JNK, p38, IKK, and I κ B α phosphorylation as well as I κ B α degradation and NF- κ B nuclear translocation. TAK1-deficient MEF cells were transduced with the retrovirus encoding the vector control, TAK1 wild type, or TAK1 K158R mutant and subsequently selected with puromycin (2 μ g/ml) to establish the TAK1-deficient MEF cell lines with the stable expression of either TAK1 wild type or K158R mutant. TAK1-deficient, TAK1 wild type, and K158R mutant reconstituted MEF cells were untreated or treated with TNF α (10 ng/ml) (A) and IL-1 β (10 ng/ml) (B) for the times indicated and subsequently harvested. Whole cell extracts and nuclear extracts were subjected to SDS-PAGE and immunoblotted (IB) with antibodies indicated. β -Actin was detected as a loading control for whole cell extracts, and PCNA was used as a loading control for nuclear extracts. C-F, expression of the TAK1 K158R mutant inhibits TNF α (C and E) and IL-1 β (D and F)-induced NF- κ B (C and D) and AP-1 (E and F) reporter activities. NF- κ B (C and D) and AP-1 (E and F) reporter and 20 ng of *Renilla*-Luc plasmids were cotransfected into TAK1-deficient MEF control, TAK1 wild type, and K158R reconstituted cells for 48 h. Cells were then either untreated or treated with TNF α (1 ng/ml) and IL-1 β (1 ng/ml) for 6 h. The relative luciferase activity was measured and normalized with the *Renilla* activity. Error bars, \pm S.D. in triplicate experiments.

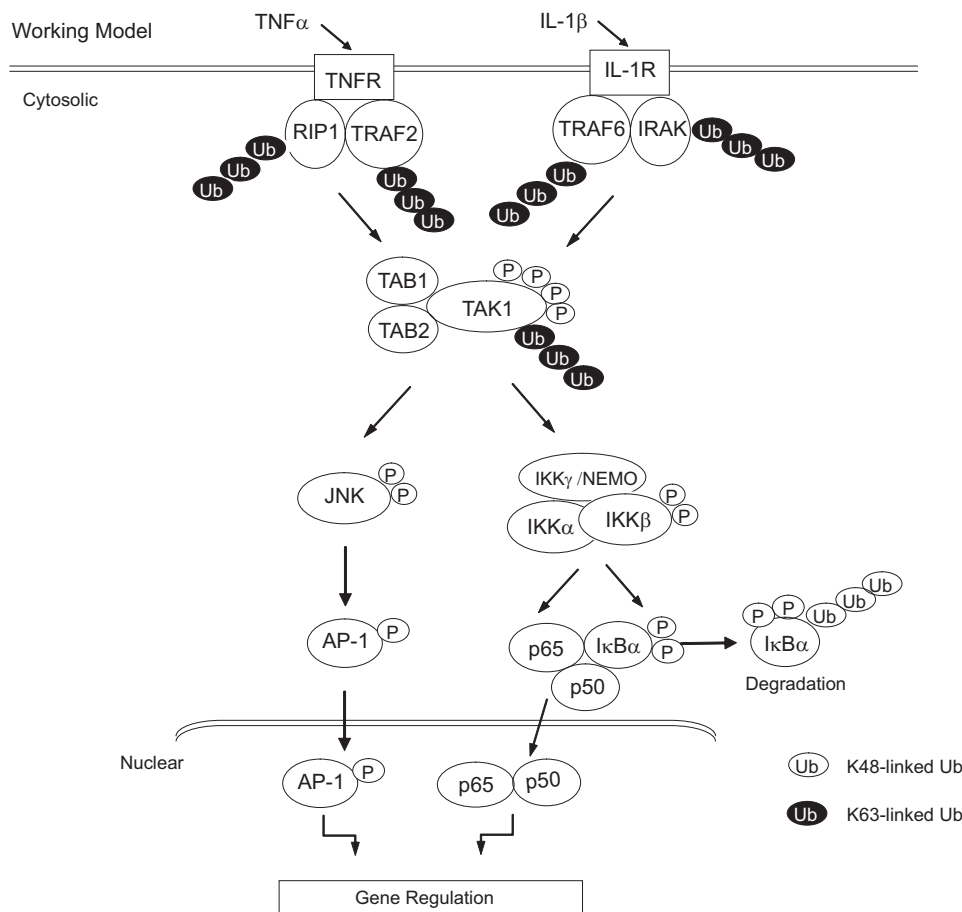


FIGURE 9. A working model for the role of Lys⁶³-linked TAK1 polyubiquitination in TNF α - and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation. TNF α and IL-1 β induces TAK1 polyubiquitination at Lys¹⁵⁸ residue and phosphorylation at the Thr¹⁷⁸, Thr¹⁸⁴, Thr¹⁸⁷, and Ser¹⁹² residues for TAK1 activation. In turn, activated TAK1 mediates IKK/NF- κ B and JNK/AP-1 activation as well as NF- κ B- and AP-1-dependent IL-6 gene expression in the cells.

of IKK γ /NEMO to the polyubiquitin chain on RIP1 and IRAKs (8–12). Our current results demonstrate that TAK1 is polyubiquitinated at the Lys¹⁵⁸ residue, and this site-specific Lys⁶³-linked polyubiquitination is required for binding of NEMO to TAK1 in response to TNF α and IL-1 β stimulation. Previous structure studies indicate that TAK1 Lys¹⁵⁸ residue is on the surface of the protein structure and exposed to solvent (50). Therefore, the TAK1 K158R mutation should not induce any significant structural perturbation (supplemental Fig. S6). Consistent with this study, we found that both TRAF2 and TRAF6 successfully catalyzed TAK1 K63R polyubiquitination in an *in vitro* ubiquitination assay (supplemental Fig. S1). Together, these results suggest that TAK1 polyubiquitination at the Lys¹⁵⁸ residue contributes to the formation of TNF α - and IL-1 β -induced intracellular signaling complex that is essential for optimal downstream IKK/NF- κ B and JNK/AP-1 activation.

Interestingly, as two serine/threonine kinases, RIP1 and IRAK1 are polyubiquitinated with a Lys⁶³ linkage type at lysine residues outside of their kinase domain in response to TNF α and IL-1 β stimulation, respectively (8–12). In addition, Lys⁶³-linked polyubiquitination of these two kinases are not reported to be involved in the regulation of their kinase activities. Furthermore, site-specific monoubiquitination of IKK β within its kinase domain regulates its persistent activation (51). In con-

trast, TNF α - and IL-1 β -induced Lys⁶³-linked TAK1 polyubiquitination within the kinase domain is required for TNF α - and IL-1 β -induced activation of TAK1 and the downstream signal transduction pathway.

Using a HeLa cell line stably expressing FLAG-TAK1 wild type and K158R mutant, we found that TNF α and IL-1 β induced binding of IKK γ /NEMO to TAK1 wild type but not K158R mutant. This result strongly indicates that TAK1 polyubiquitination at the Lys¹⁵⁸ residue is essential for TNF α - and IL-1 β -induced association of TAK1 with IKK γ /NEMO. It is highly likely that only polyubiquitinated TAK1 is able to phosphorylate and activate the IKK γ /NEMO-associated IKK β in a complex containing the ubiquitinated TAK1, IKK γ /NEMO, and IKK β . Further study is required to dissect the mechanism by which polyubiquitinated TAK1 is recruited to the receptor-associated protein complex and activates IKK/NF- κ B and JNK/AP-1 in response to TNF α and IL-1 β stimulation.

Our data show that polyubiquitination of TAK1 is not only required for TAK1-mediated IKK β activation but also for JNK and p38 activa-

tion. Importantly, we also failed to observe the phosphorylation at the Thr¹⁷⁸, Thr¹⁸⁴, and Thr¹⁸⁷ residues in the activation loop of TAK1 upon co-overexpression of TAK1 K158R with TAB1 in HEK-293T cells (data not shown). These results suggest that polyubiquitination of TAK1 at the Lys¹⁵⁸ residue is not only required for maintaining the receptor-associated complex formation but also essential for TAK1 phosphorylation and activation. Further studies are needed to determine the molecular mechanism of Lys⁶³-linked TAK1 polyubiquitination in the regulation of TAK1 activation.

During preparation of this paper, Sorrentino *et al.* (43) reported that TRAF6-mediated Lys⁶³-linked TAK1 polyubiquitination at the Lys³⁴ residue correlates with TGF- β -induced TAK1 activation. TAK1 Lys³⁴ is located at the N terminus of the TAK1 protein sequence, outside of its kinase domain. Surprisingly, we did not find any role of the Lys³⁴ residue in TRAF2- and TRAF6-mediated Lys⁶³-linked TAK1 polyubiquitination and NF- κ B activation. Therefore, more work is needed to determine whether Lys³⁴ residue is an actual Ub lysine acceptor site on TAK1 in TGF- β signaling. However, it is possible that TRAF6 mediates TAK1 polyubiquitination at different sites in different signaling pathways.

In summary, our study revealed that polyubiquitination of TAK1 at the Lys¹⁵⁸ residue within the kinase activation loop is

required for TNF α and IL-1 β -mediated IKK/NF- κ B and JNK/AP-1 activation. Importantly, our finding in this study is the first report that Lys⁶³-linked polyubiquitination of a serine/threonine kinase within the kinase activation loop is required for kinase activation and proinflammatory cytokine TNF α - and IL-1 β -mediated signal transduction. In view of the data presented here and in previous reports, we propose a working model (Fig. 9) in which, upon TNF α and IL-1 β binding to their cognate receptors, the receptor-mediated signaling events induce TRAF2/6-mediated TAK1 polyubiquitination at the Lys¹⁵⁸ and phosphorylation at the Thr¹⁷⁸/Thr¹⁸⁴/Thr¹⁸⁷/Ser¹⁹² residues within the kinase domain, which in turn trigger the activation of TAK1-mediated IKK/NF- κ B and JNK/AP-1 as well as NF- κ B- and AP-1-dependent IL-6 gene expression in the cells.

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