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Author(s)	Kashiwakura, Jun-ichi; Koizumi, Nao; Saitoh, Kodai; Kagohashi, Kota; Sasaki, Yuto; Kobayashi, Fuki; Kawahara, Shoya; Yamauchi, Yukie; Kitai, Yuichi; Muromoto, Ryuta; Oritani, Kenji; Matsuda, Tadashi
Citation	Biochemical and biophysical research communications, 572, 80-85 <a href="https://doi.org/10.1016/j.bbrc.2021.07.098">https://doi.org/10.1016/j.bbrc.2021.07.098</a>
Issue Date	2021-10-01
Doc URL	<a href="http://hdl.handle.net/2115/87011">http://hdl.handle.net/2115/87011</a>
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Type	article (author version)
File Information	Kashiwakura J BBRC.pdf



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**Signal-transducing adaptor protein-2 has a nonredundant role for IL-33-triggered mast cell activation.**

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This manuscript contains 26 pages with 3 figures.

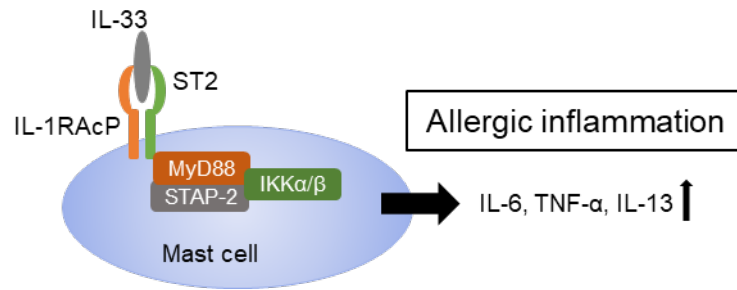
The text contains 4,118 words.

## **Abstract**

Signal-transducing adaptor protein (STAP)-2 is one of the STAP family adaptor proteins and ubiquitously expressed in a variety types of cells. Although STAP-2 is required for modification of FcεRI signal transduction in mast cells, other involvement of STAP-2 in mast cell functions is unknown, yet. In the present study, we mainly investigated functional roles of STAP-2 in IL-33-induced mast cell activation. In STAP-2-deficient, but not STAP-1-deficient, mast cells, IL-33-induced IL-6 and TNF-α production was significantly decreased compared with that of wild-type mast cells. In addition, STAP-2-deficiency greatly reduced TLR4-mediated mast cell activation and cytokine production. For the mechanisms, STAP-2 directly binds to IKKα after IL-33 stimulation, leading to elevated NF-κB activity. In conclusion, STAP-2, but not STAP-1, participates in IL-33-induced mast cells activation.

**Keywords:** Mast cells, IL-33, Signal-transducing adaptor molecule-2, ST2, NF-κB

## Graphic abstract



## Introduction

IL-33, one of the IL-1 family members, is a key cytokine for the development of Th2 immune responses. IL-33 is produced by damaged/stressed epithelial cells and cooperates with IL-25 to activate innate lymphoid cells (ILC2) [1-3]. Activated ILC2 then produce IL-5 and IL-13, resulting in driving generation of Th2 immune responses [1, 4, 5]. IL-33-deficient mice have been reported to be resistant to *Schistosoma mansoni*-induced pulmonary granuloma formation, which is dependent on a typical Th2-like immune responses [6]. The mice also show less severity of allergic immune responses, such as allergic rhinitis and airway inflammation [7]. Thus, IL-33 is likely to be a suitable target to generate a new therapeutic strategy for allergic disorders.

Mast cells (MCs) are an important cell type for IgE-mediated allergic inflammatory reactions, such as anaphylaxis. High affinity IgE receptor (FcεRI) is expressed on cell surface of MCs, and aggregation of IgE-bound FcεRI results in the secretion of granules containing histamine and proteases as well as the production of cytokines, such as IL-6 and TNF-α [8]. MCs are also activated by some stimuli except IgE, including neuropeptides, bacterial/virus components, and allergic inflammation-related cytokines [9]. IL-33 receptor, ST2, is expressed on MCs, and IL-33 accelerates mast cell cytokine production and mast cell-mediated allergic inflammatory reaction

[10-14]. IL-33 also activates human mast cells to promote maturation, survival, adhesion, and cytokine production [15, 16]. In addition, IL-33 is a key cytokine to amplify both IgE-dependent and -independent human mast cell activation [17].

Signal-transducing adaptor protein (STAP) family consists of two members, STAP-1 and STAP-2 [18-21]. STAP-2 is an important protein to regulate immune responses and tumorigenesis [22]. STAP-2, which is expressed in mast cells, negatively regulates degranulation and cytokine production through affecting FcεRI signal transduction [23]. STAP-2 also participates in TLR4 signal transduction and LPS-induced macrophage activation through direct binding to MyD88 [24], which is also an important adaptor protein for the IL-33 signal pathway [25, 26]. However, it remains unknown whether STAP-2 is involved in IL-33-induced mast cell activation and how STAP-2 affects ST2 signal transduction in mast cells. The aims of this study are to investigate effects of STAP-2 on IL-33-induced mast cell activation.

## **Materials and Methods**

### ***Mice***

Balb/c and C57BL/6 mice were purchased from SANKYO LABO SERVICE CO. Inc. (Hokkaido, Japan). Balb/c-background STAP-2-deficient (STAP-2 KO) and C57BL/6-

background STAP-1 deficient (STAP-1 KO) mice were previously generated [27, 28]. All animal studies were approved by the Hokkaido University animal ethics committee. All mice were housed and bred in the Pharmaceutical Sciences Animal Center of Hokkaido University under specific pathogen-free conditions.

### ***Antibodies***

APC anti-mouse c-Kit (clone: 2B8), FITC anti-mouse FcεRIα (clone: MAR-1), PE anti-mouse ST2 (clone: DIH4) and PE anti-mouse TLR4 (clone: MTS510) mAbs were purchased from BioLegend (San Diego, CA). Anti-phospho Akt, anti-Akt, anti-phospho IKKα/β and anti-phospho NF-κB p65 Abs were purchased from Cell Signaling Technology (Beverly, MA). Anti-Myc (clone: 9E10) and anti-FLAG (clone: M2) mAbs as well as anti-Myc polyclonal Ab were purchased from Sigma-Aldrich (St. Louis, MO). Other antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA).

### ***Generation of bone marrow-derived cultured mast cells***

Femoral bone marrow cells were cultured in 10% FCS RPMI1640 containing 3 ng/mL recombinant mouse IL-3 (TONBO biosciences, San Diego, CA, USA) for 4-6 weeks to generate bone marrow-derived cultured mast cells (BMMCs). The purity of FcεRI<sup>+</sup>c-Kit<sup>+</sup>



BMMCs was more than 90 % after culturing.

### ***Cell culture and establishment of stable transfectants***

Ba/F3 cells are cultured in RPMI1640 containing 10% FCS and 10% IL-3 condition medium. ST2-, STAP-2- and ST2/STAP-2-overexpressing Ba/F3 cells were generated using pEC-ST2-FLAG and pBabe-Myc-STAP-2 plasmids. After selection using puromycin (3 µg/mL) and G418 (1 mg/mL), drug-resistant Ba/F3 cells were cloned by limiting dilution methods. Expression FLAG-tagged ST2 and Myc-tagged STAP-2 in the clones was detected by FACS analysis and western blotting, and generated clones were used for the experiments.

### ***Flowcytometric analysis***

Flowcytometric analysis was performed as previously described [27].

### ***Cytokine production***

BMMCs were stimulated with indicated concentrations of IL-33 (TONBO biosciences) and LPS (Sigma-Aldrich) for 24 h. IL-6, TNF- $\alpha$  and IL-13 levels in supernatants were measured using ELISA kits (IL-6 and TNF- $\alpha$ ; BioLegend, IL-13; Affymetrix, San Diego,

CA).

### ***Western blotting***

Western blot analysis was performed as previously described [27]. For immunoprecipitation, cell lysates were incubated with primary Ab, followed by incubation with nProtein A Sepharose™ 4 Fast Flow (GE Healthcare Bio-Sciences) or Protein G Resin (GenScript Japan Inc., Tokyo, Japan). After beads were washed, the beads were boiled in 1x SDS sample buffer and co-immunoprecipitated proteins were detected by western blotting. Actin was detected as a loading control.

### ***Statistical analysis***

Statistical analysis was performed using GraphPad Prism 6.02. Mann-Whitney *U*-test was employed. Data were considered significant at  $p < 0.05$ . Data were shown mean + SEM.

## **Results**

### ***STAP-2, but not STAP-1, is important for IL-33-induced mast cells activation.***

We first confirmed surface expression levels of ST2 on cultured mast cells by FACS analysis. STAP-2 KO BMDCs expressed similar levels of ST2 on their surface to

WT BMMCs (Figs. 1A, B). To investigate effects of STAP-2 on IL-33-induced cytokine production, we stimulated WT and STAP-2 KO BMMCs with IL-33, and measured IL-6, TNF- $\alpha$  and IL-13 levels in supernatant. Although BMMCs produced these cytokines after IL-33 stimulation, the levels of IL-6 and TNF- $\alpha$  were significantly reduced in IL-33-stimulated STAP-2 KO BMMCs compared with WT BMMCs (Figs. 1C, D). Also, IL-13 production tended to reduce in STAP-2 KO BMMCs (Fig. 1E). Because STAP-1 is another STAP family member and involved in certain immune responses [28], we compared cytokine production of STAP-1 KO BMMCs with WT BMMCs to figure out the contribution of STAP-1 for IL-33-induced mast cell activation. Levels of IL-6, TNF- $\alpha$  and IL-13 in culture supernatant of IL-33-activated STAP-1 KO BMMCs were comparable to those of WT BMMCs (Figs 1F-H). Taken together, these results indicated that STAP-2 has an unique function for IL-33-induced mast cell activation.

***STAP-2 has nonredundant role for TLR4-mediated mast cells activation.***

IKK $\alpha$  underlies in both ST2- and TLR4-signaling pathways. Because STAP-2 binds to IKK $\alpha$  and because STAP-2 deficiency results in reduction of LPS-mediated macrophage activation [24], we evaluated the role of STAP-2 for LPS-induced mast cell activation. We first confirmed that the expression levels of TLR4 on STAP-2 KO

BMMCs were same as those on WT BMMCs (Figs. 2A, B). To investigate effects of STAP-2 on LPS-induced cytokine production, we stimulated WT and STAP-2 KO BMMCs with LPS, and measured IL-6, TNF- $\alpha$  and IL-13 levels in supernatant. The production of IL-6 was significantly reduced in STAP-2 KO BMMCs compared with WT BMMCs (Fig. 2C). The levels of TNF- $\alpha$  and IL-13 tended to reduce in STAP-2 KO BMMCs compared with WT BMMCs (Figs. 2D, E). In contrast, STAP-1 KO BMMCs produced similar levels of IL-6, TNF- $\alpha$  and IL-13 in supernatant compared with WT BMMCs (Figs 2F-H). Taken together, these results indicated that STAP-2 has a nonredundant function for TLR4-mediated cell activation in mast cells.

***STAP-2 enhances for IL-33-induced signal transduction.***

To investigate molecular mechanisms by which STAP-2 regulates IL-33 signaling, we compared ST2 signal transduction between IL-33-stimulated WT and STAP-2 KO BMMCs by western blotting. Phosphorylation levels of NF- $\kappa$ B p65, IKK $\alpha$ / $\beta$  and Akt were reduced in STAP-2 KO BMMCs compared with WT BMMCs after stimulation with IL-33 (Fig. 3A). To investigate whether STAP-2 directly regulates ST2 signal transduction or not, we prepared ST2-, STAP-2- and ST2/STAP-2-overexpressing Ba/F3 cells whose ST2 expression was confirmed by FACS analysis.

ST2 was expressed on ST2- and ST2/STAP-2-overexpressing, but not mock-transfected, Ba/F3 cells (Fig. 3B). Using these transfectants, we found that NF- $\kappa$ B signal pathway was further enhanced in ST2/STAP-2-overexpressing Ba/F3 cells compared with ST2- and STAP-2-overexpressing Ba/F3 cells after IL-33 stimulation (Fig. 3C).

We next analyzed direct interaction of STAP-2 with ST2 or IRAK4, which is involved in IL-33 signal transduction. As shown in Fig. 3D, neither ST2 nor IRAK4 were immunoprecipitated with STAP-2 in 293T cell preparations, although a positive control BRK was immunoprecipitated with STAP-2 as previously shown [29]. This result suggested that STAP-2 has no ability to directly bind to ST2 or IRAK4. We finally examined the interaction of STAP-2 with IKK $\alpha$  using ST2- and ST2/STAP-2-overexpressing Ba/F3 cells to investigate whether STAP-2 had ability to bind to IKK $\alpha$  in response to IL-33. Without the IL-33 stimulation, no association of STAP-2 with IKK $\alpha$  was observed. When the cells were stimulated with IL-33, we observed direct interaction of STAP-2 with IKK $\alpha$  in ST2/STAP-2-, but not ST2-, overexpressing Ba/F3 cells (Fig. 3E). Correctively, these results suggested that STAP-2 enhances IL-33-induced activation of NF- $\kappa$ B signal pathway by inducing interaction of STAP-2 with IKK $\alpha$  after IL-33 stimulation.

## Discussions

In this study, we described that STAP-2, but not STAP-1, is involved in IL-33-induced mast cell cytokine production. Mechanistic analysis revealed that STAP-2 interacts with IKK $\alpha$  in IL-33-stimulated cells. These results indicate that STAP-2 has a nonredundant role for ST2-mediated mast cell activation.

Mast cells are one of the important cell types for pathogenesis of allergic diseases, such as bronchial asthma and food allergies. IL-33 is critical for mast cell activation and mast cell-mediated allergic inflammation. Indeed, IL-33 induces cytokine production in mast cells, and this IL-33/mast cell/cytokine axis plays an important role for airway inflammation [11, 13, 14, 16]. Proportions of IL-33-positive cells in lungs from patients with bronchial asthma are higher than normal subjects [30]. Thus, IL-33 is believed to be a suitable therapeutic target for allergic diseases. In the present study, we demonstrated that STAP-2 deficiency results in a decrease of IL-33-induced mast cell activation. Although details of the inducing machinery are not figured out yet, STAP-2 may directly affect canonical NF- $\kappa$ B signaling because 1) phosphorylation of NF- $\kappa$ B p65 and IKK $\alpha/\beta$  is reduced in STAP-2 KO BMMCs, 2) phosphorylation of these molecules is increased in ST2/STAP-2-overexpressing Ba/F3 cells, 3) STAP-2 binds to IKK $\alpha$  in IL-33-stimulated ST2/STAP-2-overexpressing Ba/F3 cells.

STAP family consists of two members, STAP-1 and STAP-2. We previously reported that STAP-2 is required for FcεRI-mediated signal transduction and basophil activation [27]. However, our preliminary results suggest that STAP-1 is dispensable for FcεRI-mediated basophil activation (data not shown), implicating that STAP-2 has nonredundant function of FcεRI signaling in basophils. The effects on B cell receptor-mediated B cell activation are also limited to STAP-2, but not STAP-1 (In preparation). In this study, IL-33-induced mast cell activation is reduced by STAP-2-, but not STAP-1-, deficiency, suggesting that STAP-2 has a nonredundant role for IL-33 signaling in mast cells. STAP-2-restricted function is also observed when BMDCs were stimulated with LPS. Although it is unknown yet why only STAP-2 affects ST2 and TLR4 signal pathways in mast cells, a proline-rich domain within STAP-2 might be involved in its nonredundant function in mast cells because STAP-1 has no proline-rich domain [18-21]. This will be the subject of future studies.

In this report, we showed direct interaction of STAP-2 with IKKα in ST2/STAP-2-overexpressing Ba/F3 cells upon stimulation with IL-33. We have previously reported that STAP-2 phosphorylation is important for regulation of STAT3 and STAT5 activity [21, 29, 31, 32], BCR-ABL-dependent cell growth [33], its interaction with Pyk2 [34] and EGFR-mediated tumor growth [35]. Although we examined phosphorylation of

STAP-2 in IL-33-stimulated ST2/STAP-2-overexpressing Ba/F3 cells, no phosphorylation of STAP-2 in the cells was observed (data not shown), suggesting that STAP-2 phosphorylation is not required for IL-33-induced NF- $\kappa$ B signal transduction.

In summary, our study demonstrated that STAP-2 deficiency results in reduction of cytokine production from mast cells upon IL-33 stimulation. For the mechanisms, we propose that STAP-2 interacts with IKK $\alpha$  after IL-33 stimulation. Thus, we suggest that STAP-2 is essential for promoting IL-33-induced mast cell activation and allergic inflammation.

### **Acknowledgements**

We appreciate Drs. Masaaki Hashiguchi (Dokkyo Medical University) and Shintaro Sato (Osaka University) for providing the ST2-FLAG and FLAG-IRAK vectors, respectively.

We thank all members of Hokkaido University, Global Facility Center (GFC), Pharma Science Open Unit (PSOU) funded by MEXT under “Support Program for Implementation of New Equipment Sharing System”.

### **Funding**

This study was partly supported by JSPS KAKENHI Grants 19H03364 (T.M.) and



21K08451 (J.K.).

### **Conflict of Interest (COI)**

The authors declare no conflict of interest.

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## Figure Legends

**Fig. 1.** Cytokine production in STAP-2 KO mast cells upon stimulation with IL-33.

(A) Expression of ST2 on WT and STAP-2 KO BMMCs. Gray-filled and black line histograms are unstained and stained BMMCs, respectively. (B) Mean fluorescence intensity of the ST2 expression on WT and STAP-2 KO BMMCs. White and black bars are WT and STAP-2 KO BMMCs, respectively. Data shown are mean + SEM of 4 independent experiments (WT = 4, KO = 6). (C-E) Levels of IL-6 (C), TNF- $\alpha$  (D) and IL-13 (E) in supernatants of WT and STAP-2 KO BMMCs after stimulation with IL-33



for 24 h. White and black bars are WT and STAP-2 KO BMMCs, respectively. Data shown are mean + SEM of 5 (IL-6), 4 (TNF- $\alpha$ ) and 6 (IL-13) independent experiments (IL-6; n = 10, TNF- $\alpha$ ; n = 8, IL-13; n = 11). (F-H) Levels of IL-6 (F), TNF- $\alpha$  (G) and IL-13 (H) in supernatants of WT and STAP-1 KO BMMCs after stimulation with IL-33 for 24 h. White and gray bars are WT and STAP-1 KO BMMCs, respectively. Data shown are mean + SEM of 3 (IL-6) and 2 (TNF- $\alpha$  & IL-13) independent experiments (IL-6; n = 6, TNF- $\alpha$ ; n = 4, IL-13; n = 3). \*, p<0.05, \*\*, p<0.01 by Mann-Whitney *U*-test. ns = no significance

**Fig. 2.** Cytokine production in STAP-2 KO mast cells after LPS stimulation.

(A) Expression of TLR4 on WT and STAP-2 KO BMMCs. Gray-filled and black line histograms are unstained and stained BMMCs, respectively. (B) Mean fluorescence intensity of the LPS expression on WT and STAP-2 KO BMMCs. White and black bars are WT and STAP-2 KO BMMCs, respectively. Data shown are mean + SEM of 3 independent experiments (n = 3). (C-E) Levels of IL-6 (C), TNF- $\alpha$  (D) and IL-13 (E) in supernatants of WT and STAP-2 KO BMMCs after stimulation with LPS for 24 h. White and black bars are WT and STAP-2 KO BMMCs, respectively. Data shown are mean + SEM of 4 (IL-6 & TNF- $\alpha$ ) and 5 (IL-13) independent experiments (IL-6; n = 6, TNF- $\alpha$ ;

n = 6, IL-13; n = 7). (F-H) Levels of IL-6 (F), TNF- $\alpha$  (G) and IL-13 (H) in supernatants of WT and STAP-1 KO BMMCs after stimulation with LPS for 24 h. White and gray bars are WT and STAP-1 KO BMMCs, respectively. Data shown are mean + SEM of 4 (IL-6 & TNF- $\alpha$ ) and 3 (IL-13) independent experiments (IL-6; n = 8, TNF- $\alpha$ ; n = 8, IL-13; n = 6). \*, p<0.05, \*\*, p<0.01 by Mann-Whitney *U*-test. ns = no significance

**Fig. 3.** STAP-2 enhances NF- $\kappa$ B signaling by direct binding to IKK $\alpha$  in response to IL-33

(A) Comparison of activation of NF- $\kappa$ B and Akt signaling between WT and STAP-2 KO BMMCs upon stimulation with IL-33. (B) Expression of ST2 on transfected Ba/F3 cells. Gray-filled, black normal line and black thick line histograms are mock-transfected, ST2- and ST2/STAP-2-overexpressing Ba/F3 cells, respectively. (C) Comparison of activation of NF- $\kappa$ B pathway among mock-transfected, ST2-, STAP-2- and ST2/STAP-2-overexpressing Ba/F3 cells after IL-33 activation. (D) Myc-tagged STAP-2 vector was cotransfected with either FLAG-tagged IRAK4, ST2 or BRK vector into 293T cells. Binding of STAP-2 to FLAG-tagged proteins was analyzed by immunoprecipitation, followed by western blotting. (E) Interaction of Myc-tagged STAP-2 with endogenous IKK $\alpha$  in ST2- and ST2/STAP-2-overexpressing Ba/F3 cells in the presence or absence of

IL-33 stimulation. Arrowhead shows coimmunoprecipitated IKK $\alpha$ . All data shown are representative of 2-3 independent experiments.

Figure 1

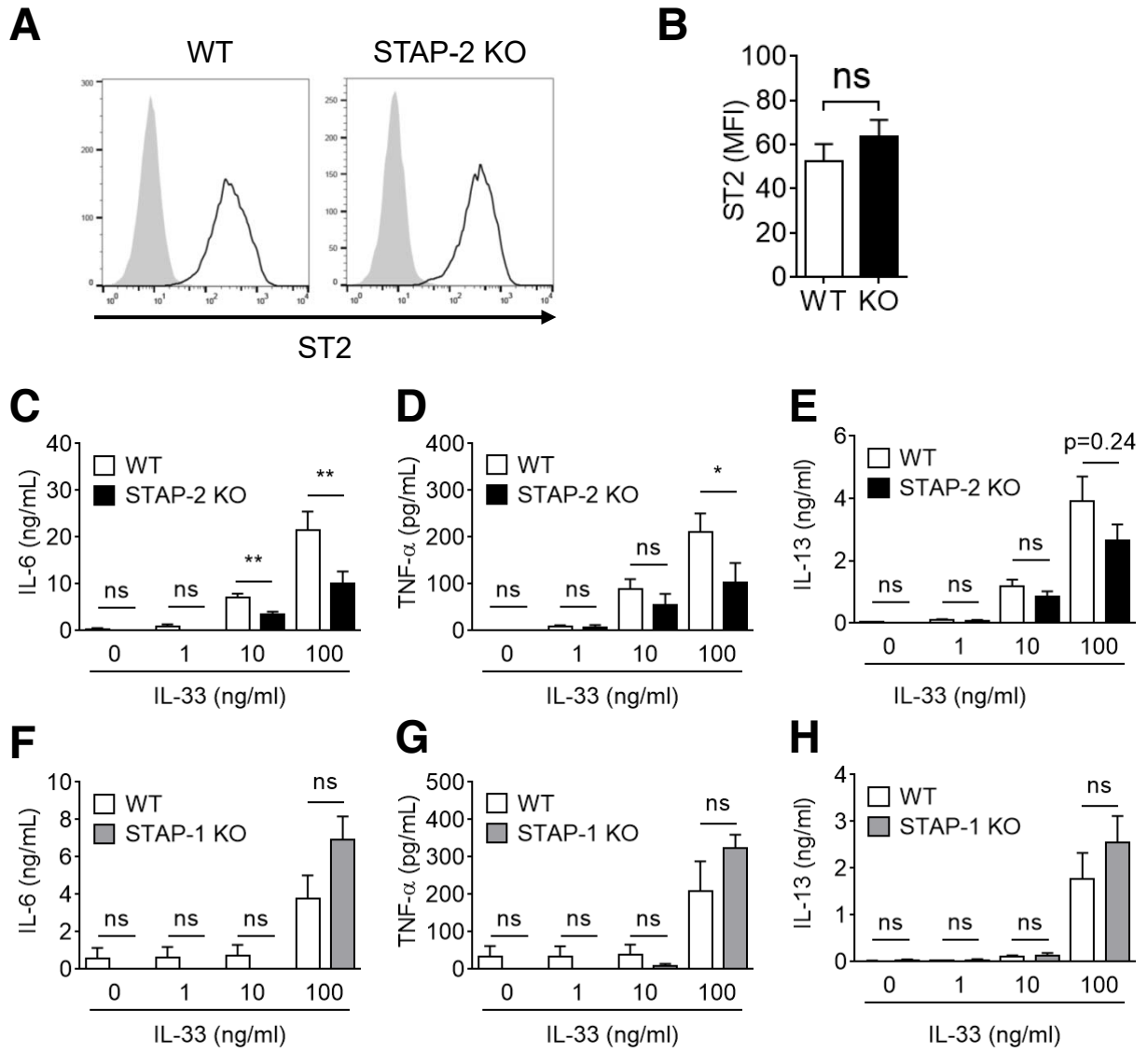


Figure 2

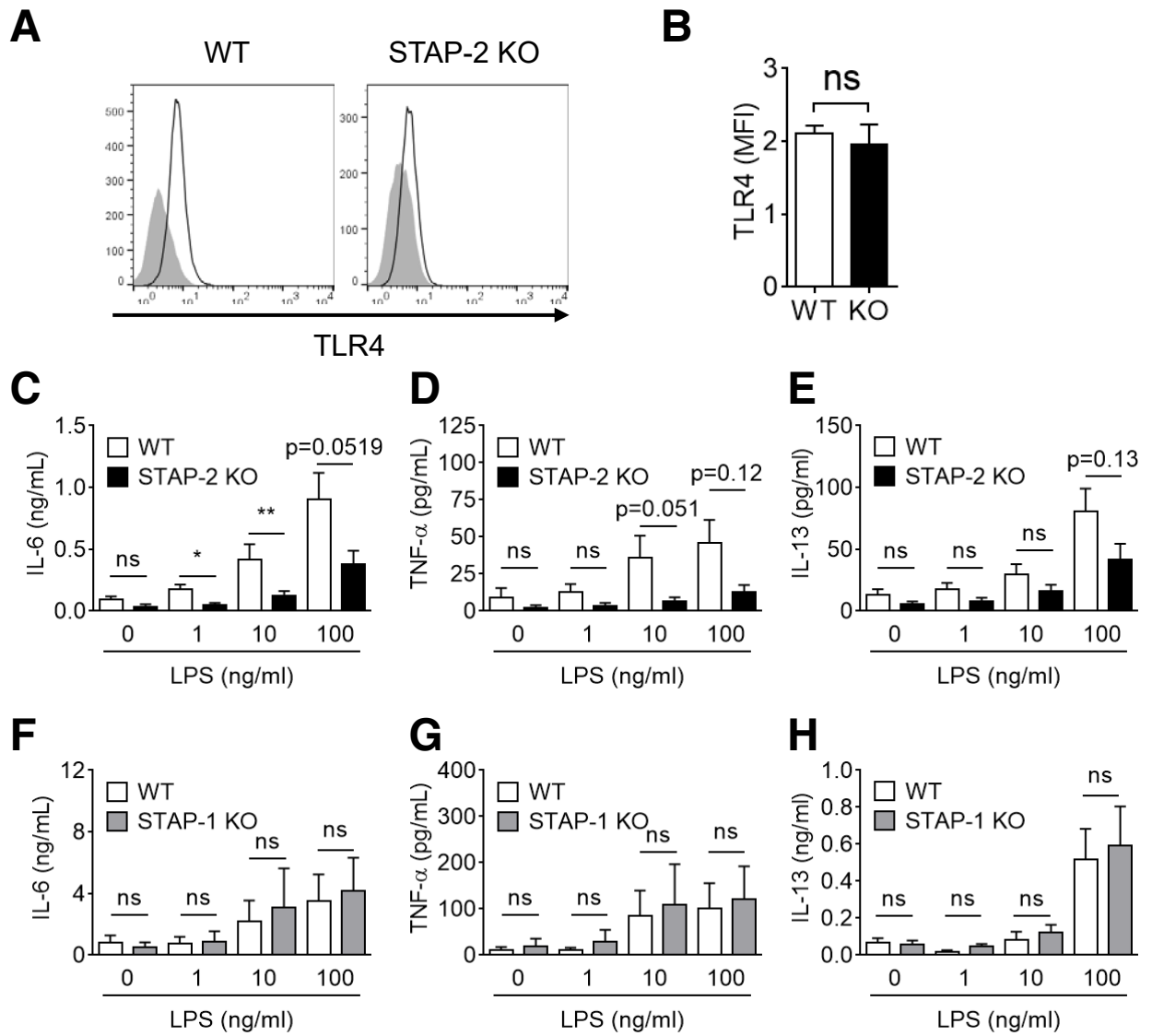


Figure 3

