



## PARP-1 Negatively Regulates Treg Cell Function

### Experimental Procedures

**Plasmids, Antibodies, and Reagents**—pIP-HA/Myc/FLAG-tagged FOXP3, FOXP3 truncations, His-tagged Ub, and FLAG-tagged Stub1 were constructed as described previously (20). PARP-1 was amplified from human peripheral blood mononuclear cell (PBMC) cDNA with the primers 5'-ATCATGGCG-GAGTCTTCGGATAAGC-3' (forward) and 5'-CCGCTC-GAGTTACCACAGGGAGGTCTTAAAATTG-3' (reverse). The antibodies used were as follows: anti-FLAG (catalog no. M2, Sigma), anti-HA (catalog no. F-7, Santa Cruz Biotechnology), anti-Myc (Santa Cruz Biotechnology), anti-FOXP3 (catalog no. hFOXY, eBioscience), anti-PARP-1 (Sigma), anti-PAR (catalog no. 4335-AMC-050, Trevigen), anti-GAPDH (catalog no. 1C4, Sungene Biotech), anti- $\alpha$ -tubulin (catalog no. DM1A, Sigma), anti-ubiquitin (Santa Cruz Biotechnology), anti-Lamin B (Santa Cruz Biotechnology), anti-H3 (catalog no. 9715, Cell Signaling Technology), anti- $\beta$ -actin (catalog no. 6G3, Sungene Biotech), anti-CD25-PE (catalog no. BC96, Biolegend), anti-CD4-FITC (catalog no. RPA-T4, Biolegend), and anti-CD127-PE-cy7 (catalog no. eBioRDR5, eBioscience). The PARP-1-specific inhibitor AG14361 (catalog no. S2178) and the PARP inhibitor 3-aminobenzamide (3-AB) were purchased from Selleck and Santa Cruz Biotechnology, respectively. MG132 was purchased from Merck (catalog no. 474790).

**Cells and Transfection**—HEK293T cells were cultured in DMEM supplemented with 10% FCS and 1% penicillin/streptomycin. Human Jurkat T cells were grown in RPMI 1640 medium supplemented with 10% FCS, 1% penicillin/streptomycin, 1% sodium pyruvate, and 1% non-essential amino acids. All cells were maintained at 37 °C in a 5% CO<sub>2</sub> incubator. HEK293T cells were transfected with the indicated plasmids using polyethylenimine (Polyscience, catalog no. 23966-2) reagent according to the instructions of the manufacturer.

**Isolation of Human Treg Cells and Naive T Cells**—Human PBMCs were isolated from the buffy coat of healthy donors (Shanghai Blood Center). Human CD4<sup>+</sup>CD25<sup>hi</sup>CD127<sup>lo</sup> Treg cells and CD4<sup>+</sup>CD127<sup>hi</sup>CD45RA<sup>hi</sup> naive T cells were sorted using a FACS ARIA II cell sorter (BD Biosciences). The purity of the isolated cells was 95–99%. *In vitro* expansion of Treg cells was performed in X-VIVO (Lonza) medium supplemented with 10% human AB serum, 1% GlutaMax (Gibco), 1% sodium pyruvate (Gibco), and 500 units/ml rIL-2 (R&D Systems) in the presence of anti-human CD3/CD28-conjugated Dynabeads (Invitrogen) at a bead-to-cell ratio of 4:1. The purified naive T cells were differentiated into iTreg (induced Treg) cells with 5 ng/ml TGF- $\beta$  (R&D Systems), 100 units/ml rIL-2, and anti-human CD3/CD28-conjugated Dynabeads at a ratio of 1:1.

**Quantitative Real-time PCR**—Total RNA was isolated from whole PBMCs using TRIzol reagent (Invitrogen) and following the instructions of the manufacturer. RNA was quantified, and cDNA was reverse-transcribed with the PrimeScript RT reagent kit (TakaRa). The cDNA samples were used at 10 ng/well in a 384-well plate and run in triplicate. PCR reactions were set up in 10- $\mu$ l volumes with SYBR Premix Ex Taq reagent (TakaRa) on an ABI 7900HT sequence detection system. Quantification of the target mRNA expression level was normalized to  $\beta$ -actin expression. The RT primers of human genes used

were as follows: *PARP-1* forward, 5'-AAGCCCTAAAGGCT-CAGAACG-3'; *PARP-1* reverse, 5'-ACCATGCCATCAGC-TACTCGGT-3'; *CD25* forward, 5'-GAGACGTCCATATT-TACAACAG-3'; *CD25* reverse, 5'-CCTTTGATTTCACTTGGCTTC-3'; *CTLA4* forward, 5'-CTTCTCTTCATCCCTGTCTTC-3'; *CTLA4* reverse, 5'-AAGGTCAACTCATTCCCC-ATC-3'; *FOXP3* forward, 5'-TCCCAGAGTTCCCTCCACAAC-3'; *FOXP3* reverse, 5'-ATTGAGTGTCCGCTGCTTCT-3'; *Il10* forward, 5'-TGCAAAACCAAACCACAAGA-3'; *Il10* reverse, 5'-TCTCGGAGATCTCGAAGCAT-3';  $\beta$ -actin forward, 5'-GGACTTCGAGCAAGAGATGG-3'; and  $\beta$ -actin reverse, 5'-AGCACTGTGTTGGCGTACAG-3'.

**Immunoprecipitation and Immunoblot Analysis**—Cells were lysed in radioimmune precipitation assay buffer containing 50 mM Tris/HCl (pH 7.4), 0.5% Nonidet P-40, 0.25% sodium deoxycholate, 150 mM NaCl, 1 mM EDTA with 1 mM PMSF, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 1 mM NaF, and protease inhibitor (Sigma). The supernatants were immunoprecipitated with 1  $\mu$ g of the indicated antibodies and 10  $\mu$ l of protein A/G Plus-agarose (Santa Cruz Biotechnology), followed by separation in SDS/PAGE and analysis with Western blotting.

**Ubiquitin Pulldown Assay**—After 4-h treatment with 5  $\mu$ M MG132, cells were washed with ice-cold 1 $\times$  PBS and lysed in urea buffer (10 mM Tris/HCl (pH 8.0), 8 M urea, 100 mM Na<sub>2</sub>HPO<sub>4</sub>, 0.2% Triton X-100, 1 mM *N*-ethylmaleimide, and 10 mM imidazole) for 30 min. The lysates were incubated with nickel-nitrilotriacetic acid beads (catalog no. 30210, Qiagen) for 3 h at room temperature. The beads were washed three times with urea buffer (pH 8.0) before incubation. After 3 h of incubation, the beads were washed twice with urea buffer (pH 8.0) buffer, twice with urea buffer (pH 6.3) (10 mM Tris/HCl (pH 6.3), 8 M urea, 100 mM Na<sub>2</sub>HPO<sub>4</sub>, 0.2% Triton X-100, and 10 mM imidazole), and once with a wash buffer (20 mM Tris/HCl (pH 8.0), 100 mM NaCl, 20% glycerol, 1 mM dithiothreitol, and 10 mM imidazole). Ubiquitination levels were evaluated by Western blotting with specific antibodies as indicated.

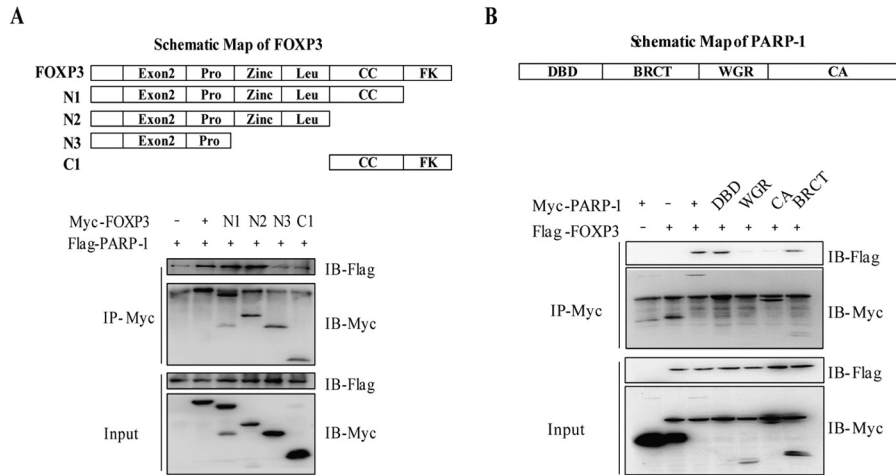
**Cytoplasm, Nucleus, and Chromatin Extraction**—1  $\times$  10<sup>6</sup> cells were harvested in 1 ml of ice-cold 1 $\times$  PBS buffer and then suspended in 300  $\mu$ l of cytoplasm buffer (10 mM Tris/HCl (pH 7.5), 10 mM KCl, 0.1 mM EDTA, 1 mM DTT, and 0.5 mM PMSF). Nonidet P-40 was added to a final concentration of 0.6% after 15 min incubation on ice. Another 15 min later, the lysate was centrifuged, and the supernatant was cytoplasm fraction. The pellet was resuspended in 200  $\mu$ l of nucleus extract buffer (20 mM Tris/HCl (pH 8.0), 400 mM NaCl, 1 mM EDTA, 1 mM DTT, 1 mM PMSF, 1 mM Na<sub>3</sub>VO<sub>4</sub>, and 1 mM NaF) and then incubated on ice for another 30 min. After centrifugation, the supernatant was nucleus fraction, and the pellet was chromatin fraction.

**In Vitro Poly(ADP-ribosylation Assay**—An *in vitro* poly(ADP-ribosylation) assay was performed as described previously (23). MBP-FOXP3 protein was expressed in *Escherichia coli* and purified with amylose resin as described previously (20). His-PARP-1 recombinant protein was bought from Sino. Purified MBP-FOXP3 (1  $\mu$ g) and His-PARP-1 (100 ng) were incubated at 37 °C with NAD (50  $\mu$ M), DTT (1 mM), MgCl<sub>2</sub> (10 mM), Tris/HCl (pH 7.4) (100 mM), and 10 mg of braked salmon germ DNA for 1 h. The reaction was stopped with 2 $\times$  loading buffer followed by Western blotting, and anti-PAR antibody

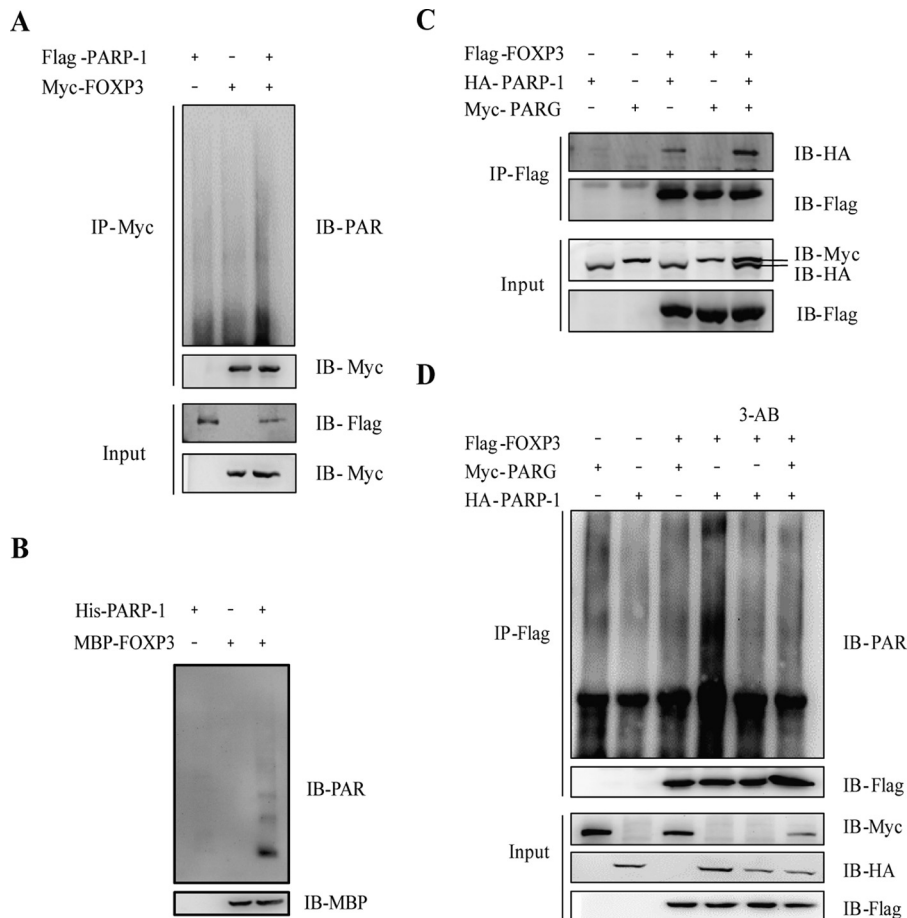




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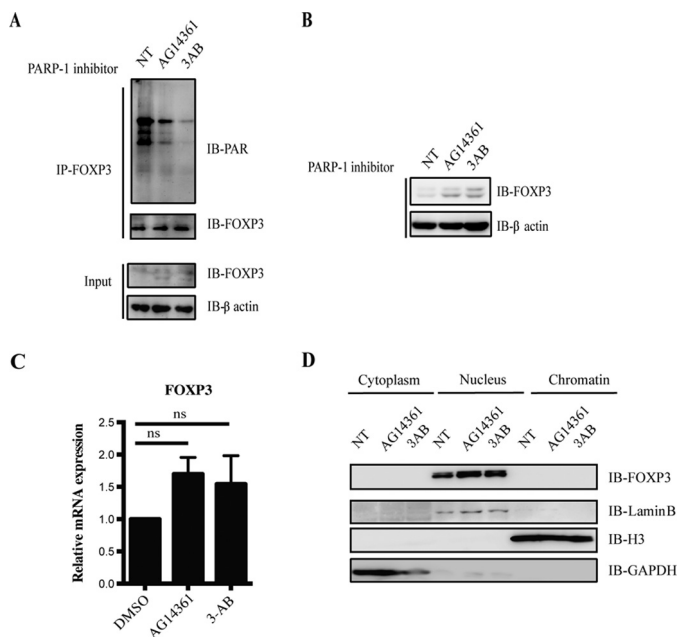
**FIGURE 2. The zinc finger/Leu zipper domains of FOXP3 interact with the DNA-binding domain/BRCA1 C terminus of PARP-1.** A, schematic of the FOXP3 protein. HEK293T cells were transfected with plasmids expressing various truncated FOXP3 proteins, followed by immunoprecipitation (IP) and Western blotting. IB, immunoblot; CC, coiled-coil domain; FK, forkhead domain. B, different domains of PARP-1 were co-expressed with FOXP3 in HEK293T cells, and the interaction of the PARP-1 domain with FOXP3 was examined by immunoprecipitation and immunoblotting. Data represent at least two independent experiments. DBD, DNA-binding domain; BRCT, BRCA1 C terminus; WGR, tryptophan-glycine-arginine domain; CA, catalytic domain.



**FIGURE 3. PARP-1 poly(ADP)-ribosylates FOXP3.** A, immunoprecipitation (IP) of exogenous Myc-FOXP3 from HEK293 T cells that were co-transfected with Myc-FOXP3 and FLAG-PARP-1 plasmids. PARylation was detected by anti-PAR antibody. IB, immunoblot. B, poly(ADP-ribosylation) assay *in vitro*. Recombinant His-tagged PARP-1 was incubated with MBP-tagged FOXP3 in poly(ADP-ribosylation) reaction buffer, and anti-PAR antibody was used to detect PARylated FOXP3. C, HEK293T cells were co-transfected with PARP-1, PARG, and FOXP3. After harvesting, immunoprecipitation and Western blotting were performed. D, HEK293T cells were co-transfected with plasmids as indicated, and then 10 mM 3-AB, a PARP inhibitor, was added to cell culture and incubated for 6 h. Cell lysates were immunoprecipitated with anti-FLAG antibody, and PARylated FOXP3 proteins were detected by anti-PAR antibody. Data represent at least two independent experiments.

lular localization, and transcriptional activity (4). To further investigate how PARP-1 inhibitors regulate FOXP3 function by inhibiting its poly(ADP-ribosylation), we first set out to exam-

ine FOXP3 protein and mRNA levels in Treg cells treated with 3-AB or AG14361. We found that, 12 h after human primary Treg cells had been treated with 3-AB or AG14361, the protein

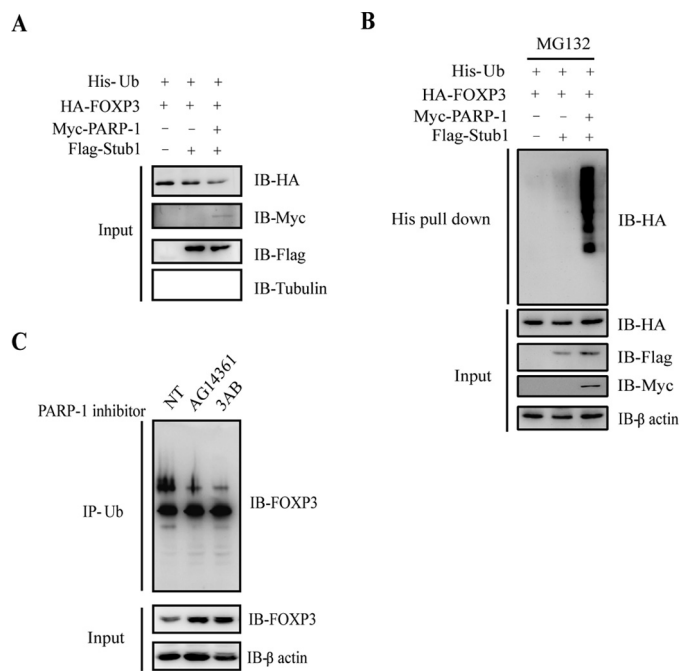


**FIGURE 4. PARP-1 inhibitors stabilize the FOXP3 protein level but not cellular localization in human Treg cells.** *A*, human primary Treg cells were cultured with the PARP-1 inhibitors 3-AB (10 mM) or AG14361 (10  $\mu$ M) for 12 h. Cell lysates were immunoprecipitated (IP) with anti-FOXP3 antibody, and PARylated FOXP3 proteins were detected by anti-PAR antibody. *IB*, immunoblot; *NT*, no treatment. *B* and *C*, human primary Treg cells were cultured with the PARP-1 inhibitors 3-AB (10 mM) or AG14361 (10  $\mu$ M) for 12 h. The protein level of FOXP3 was detected by Western blotting, and mRNA level of FOXP3 was examined by RT-PCR. *DMSO*, dimethyl sulfoxide; *ns*, no significance (two-paired Student's *t* test). Data represent at least three independent experiments. *D*, human primary Treg cells were cultured with the PARP-1 inhibitors 3-AB (10 mM) or AG14361 (10  $\mu$ M) for 12 h. Cytoplasm, nucleus, and chromatin fractions were extracted followed the protocol detailed under "Experimental Procedures" and detected by Western blotting. Data represent at least two independent experiments.

level of FOXP3 was up-regulated, but the mRNA level did not change (Fig. 4, *B* and *C*), indicating that PARP-1 inhibitors stabilized FOXP3 protein level in Treg cells through inhibiting poly(ADP-ribosyl)ation of FOXP3.

Also, we detected the cellular localization of FOXP3 in PARP-1 inhibitor-treated Treg cells and found increased FOXP3 after PARP-1 inhibitor treatment. However, FOXP3 localization still did not change (Fig. 4*D*), suggesting that the suppression of PARP-1 enzymatic activity stabilized FOXP3 but not localization in human Treg cells.

**PARP-1 Promotes the Poly-ubiquitination and Degradation of FOXP3**—The E3 ubiquitin ligase Stub1 has already been identified to negatively regulate the suppressive function of Treg cells by promoting the poly-ubiquitination and degradation of FOXP3 (20). Given the fact that the inhibition of PARP-1 enzymatic activity not only inhibited the poly(ADP-ribosyl)ation of FOXP3 but also stabilized FOXP3 in Treg cells, we decided to further examine whether PARP-1 promoted the poly-ubiquitination and degradation of FOXP3. First, His-tagged Ub, HA-tagged FOXP3, Myc-tagged PARP-1, and FLAG-tagged Stub1 were co-transfected into HEK293T cells. As expected, the FOXP3 level was reduced with addition of Stub1, and, importantly, even less FOXP3 was found in the Stub1, PARP-1, and FOXP3 co-expressed group, suggesting that PARP-1 may promote FOXP3 degradation mediated by



**FIGURE 5. PARP-1 promotes the poly-ubiquitination and degradation of FOXP3.** *A*, HEK 293T cells were co-transfected with plasmids encoding for PARP-1, FOXP3, Ub, and Stub1. Cell lysates were examined by immunoblotting (IB) with different antibodies as indicated. *B*, HEK 293T cells were co-transfected with plasmids encoding for PARP-1, FOXP3, Ub, and Stub1. Before harvesting, these cells were treated with 5  $\mu$ M MG132 for 4 h. The ubiquitin pulldown assay for His-tagged Ub was performed as described under "Experimental Procedures." *C*, human primary Treg cells were immunoprecipitated (IP) with anti-Ub antibody and analyzed by Western blotting. Data represent at least two independent experiments.

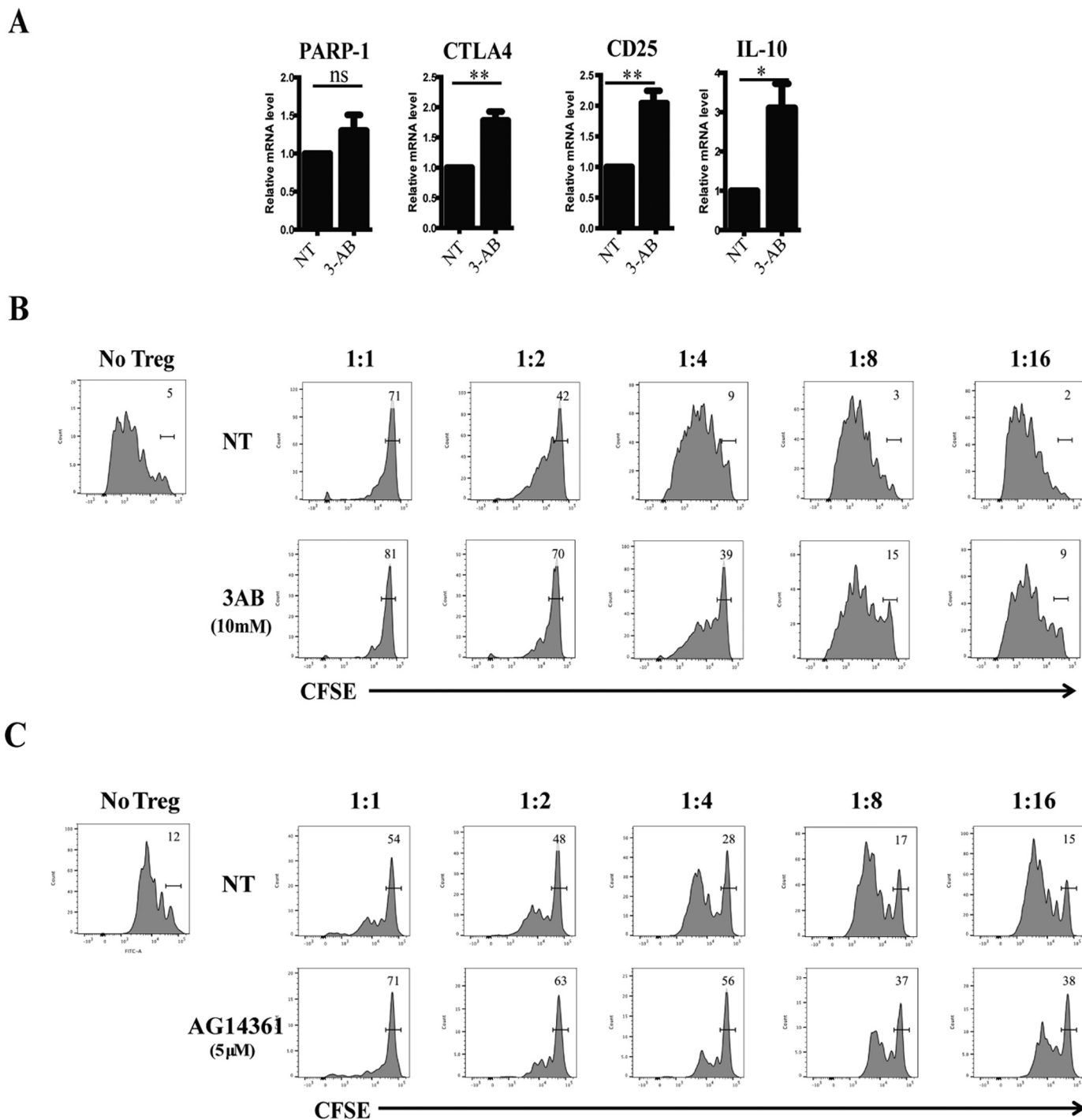
Stub1 (Fig. 5*A*). To further examine whether PARP-1 affected FOXP3 poly-ubiquitination to promote FOXP3 degradation, a His pulldown assay was used to detect the endogenous poly-ubiquitination level of FOXP3 after 4 h of MG132 treatment. The addition of the proteasome inhibitor MG132 prevented FOXP3 loss, suggesting that this process was proteasome-dependent. Moreover, the poly-ubiquitination level of FOXP3 was up-regulated significantly by addition of PARP-1, indicating that PARP-1 promoted FOXP3 poly-ubiquitination and degradation (Fig. 5*B*).

To determine whether the enzymatic activity of PARP-1 affected FOXP3 poly-ubiquitination, we detected the endogenous poly-ubiquitination level of FOXP3 in PARP-1 inhibitor-treated Treg cells. After 12 h of treatment, we found decreased FOXP3 in Ub immunoprecipitates after PARP-1 inhibitor treatment (Fig. 5*C*), suggesting that the inhibition of PARP-1 enzymatic activity suppressed the poly-ubiquitination and degradation of FOXP3.

**PARP-1 Inhibitors Enhanced the Suppressive Function of Treg Cells through PARylated FOXP3**—To further investigate whether PARP-1 inhibitors regulated Treg cell function by inhibiting FOXP3 poly(ADP-ribosyl)ation, we examined some of the FOXP3-associated genes in Treg cells treated with 3-AB. We found that, 6 h after human primary Treg cells had been treated with 3-AB, the expression levels of *CD25*, *CTLA4*, and *IL10* in the cells were up-regulated (Fig. 6*A*), suggesting that PARP-1 inhibitor affected FOXP3 downstream genes. Given the fact that the inhibition of PARP-1 enzymatic activity not



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**FIGURE 6. PARP-1 inhibitors regulate Treg cell suppressive function through inhibiting PARylated FOXP3.** *A*, human primary Treg cells were cultured with the PARP-1 inhibitor 3-AB for 6 h. The mRNA levels of FOXP3 downstream genes in selected cells were examined by RT-PCR. \*,  $p < 0.05$ ; \*\*,  $p < 0.02$ ; ns, not significant (two-paired Student's *t* test); NT, no treatment. Data represent at least three independent experiments. *B*, *in vitro* suppression assay. Human primary Treg cells were treated with or without 10 mM PARP inhibitor 3-AB for 6 h prior to the suppression assay. The Treg cells were then cultured with CFSE-labeled human PBMCs and anti-CD3/CD28 beads at the indicated ratio for 4 days. Proliferation of CD8<sup>+</sup> T effector cells was examined by flow cytometry. *C*, human primary Treg cells were treated with or without 5 µM PARP-1 inhibitor AG14361 for 12 h. Then Treg cells were cultured with CFSE-labeled human PBMCs and anti-CD3/CD28 beads at the indicated ratio for 4 days in the *in vitro* suppression assay. The proliferation of CD8<sup>+</sup> T effector cells was examined by flow cytometry. Data represent at least two independent experiments.

only inhibited the poly(ADP-ribosyl)ation of FOXP3 but also regulated FOXP3 stabilization and its downstream genes, we decided to further examine the suppressive function of Treg cells in the standard Treg suppression cell co-culture system *in vitro*. For some of the cell culture groups, Treg cells were pre-

treated with the PARP-1 inhibitor 3-AB for 6 h before the suppression assay. Human PBMCs from healthy donors were labeled with CFSE before being cultured with human primary Treg cells at different ratios and stimulated with anti-CD3/CD28 beads for 4 days. The results showed that 3-AB-pre-

treated Treg cells exhibited a much stronger suppressive effect than untreated Treg cells on the proliferation of PBMCs (Fig. 6B), suggesting that the poly(ADP-ribosylation) of FOXP3 mediated by PARP-1 negatively regulates the suppressive function of Treg cells. We also used another PARP-1-specific inhibitor, AG14361, and found that AG14361 pretreated Treg cells had an increased suppressive function *in vitro* as well (Fig. 6C). All of these data strongly suggest that PARP-1 inhibitors promote the suppressive function of Treg cells through the inhibition of FOXP3 poly(ADP-ribosylation).

## Discussion

In this study, we demonstrated that PARP-1 interacted with human FOXP3 and promoted its poly(ADP-ribosylation). We identified the zinc finger/Leu zipper domains of FOXP3 and the DNA-binding domain/BRCA1 C terminus of PARP-1 as crucial domains for the direct interaction between these two proteins. We also showed that the poly(ADP-ribosylation) of FOXP3 mediated by PARP-1 was inhibited by PARG or PARP-1 inhibitors. Most importantly, we found that reduced poly(ADP-ribosylation) of FOXP3 by PARP-1 inhibitor stabilized FOXP3, up-regulated the expression levels of FOXP3 downstream genes in Treg cells, and enhanced the suppressive function of Treg cells. Our results suggest that the poly(ADP-ribosylation) of FOXP3 mediated by PARP-1 negatively regulates the suppressive function of Treg cells.

Studies have reported that PARylation does not act alone but in concert with other posttranslational modifications (4). We found PARP-1 promoted FOXP3 poly-ubiquitination and degradation mediated by Stub1. In addition, the inhibition of PARP-1 enzymatic activity suppressed the poly-ubiquitination and degradation of FOXP3, suggesting that poly(ADP-ribosylation) of FOXP3 mediated by PARP-1 affected FOXP3 poly-ubiquitination and degradation. However, the relationship and mechanism between these two important posttranslational modifications still needs to be investigated further.

PARP-1 is an abundant nuclear protein with low enzyme activity that can be activated by DNA break, reactive oxygen species, and inflammation. The activation of PARP-1 results in the poly(ADP-ribosylation) of its target proteins, which leads to the modulation of target protein transcriptional activity, localization, and the creation of new protein interaction scaffolds (25). Our data show that PARP-1 regulates the suppressive function of Treg cells through FOXP3 poly(ADP-ribosylation). However, the regulation of PARP-1 activity and the signal induces FOXP3 poly(ADP-ribosylation) in Treg cells are still unclear. Besides, PARP-1 takes part in several cellular processes, including cell death, transcriptional regulation, and inflammation. Additional studies will be needed to figure out how these diverse functions of PARP-1 are integrated and controlled in different cells.

Because of its diverse and important roles, a number of PARP-1 inhibitors are under clinical development for the treatment of cancer, such as iniparib (BSI-201), Olaparib (AZD-2281, oral) and veliparib (ABT-888, oral) (11, 12). Our study not only has implications in the developing PARP-1 inhibitors as potential agents for the treatment of autoimmune diseases but also shows the potential risk for cancer treatment.

In summary, here we uncovered the previously unrecognized molecular mechanism that PARP-1 regulated the suppressive function of Treg cells at a posttranslational modification level through poly(ADP-ribosylation) of FOXP3. Furthermore, more specific PARP-1 inhibitors will be required both as tools and therapeutics of autoimmune diseases on the basis of the role of PARP-1 in the immune system.

**Author Contributions**—X. L. designed, performed, and analyzed the experiments shown in Figs. 1–3 and 6. J. N. and S. W. designed, performed, and analyzed the experiments shown in Figs. 4 and 5. Z. C. purified the MBP-FOXP3 protein and constructed vectors for FOXP3 truncations. W. J. C., D. L., and H. H. provided technical assistance and helped with the preparation of the manuscript. B. L. conceived and coordinated the study, analyzed and interpreted data, and wrote the paper. All authors reviewed the results and approved the final version of the manuscript.

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**Immunology:**

**Poly(ADP-ribosylation) of FOXP3 Protein Mediated by PARP-1 Protein Regulates the Function of Regulatory T Cells**

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