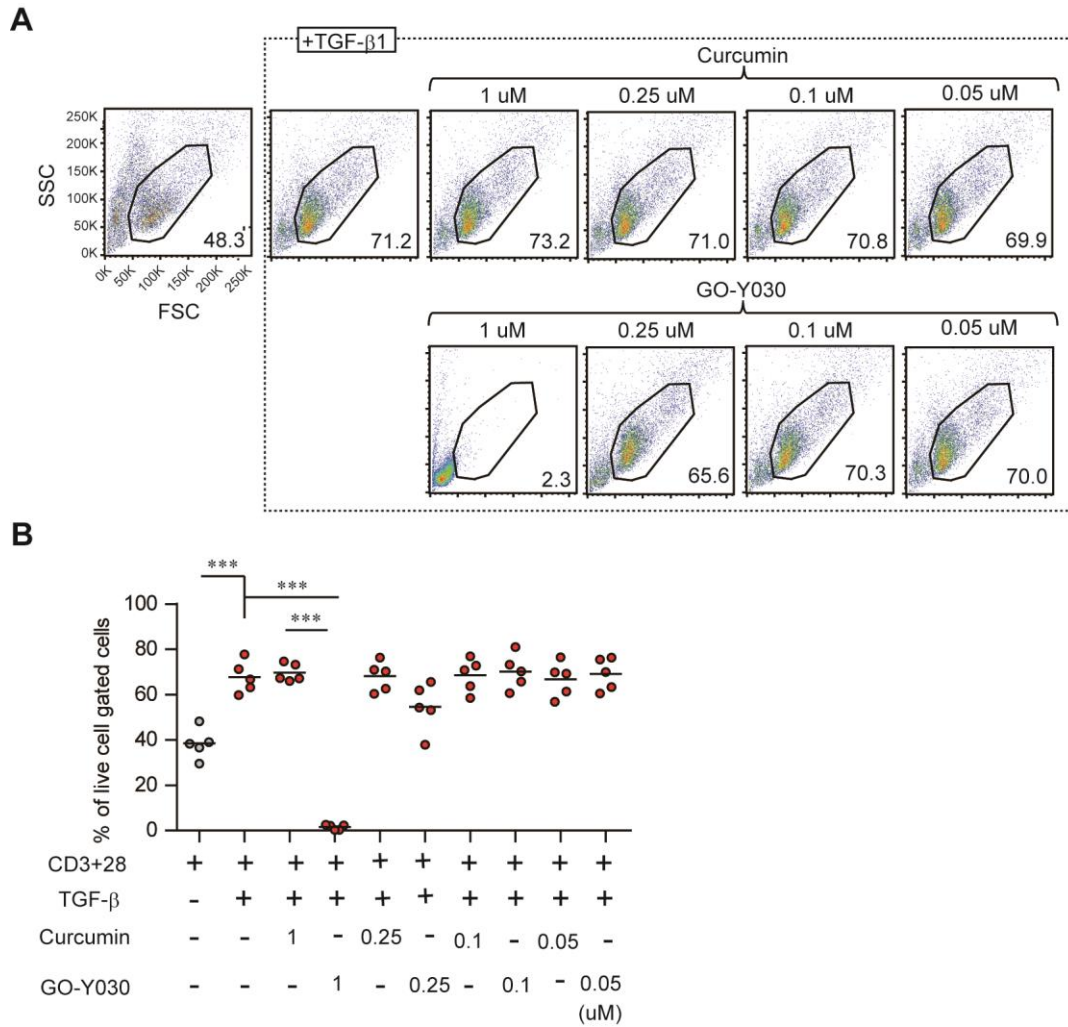


Supplementary Material

Interleukin-10	Forward	5'-CCCATTCTCGTCACGATCTC-3'
	Reverse	5'-TCAGACTGGTTTGGGATAGGTTT-3'
TGF- $\beta$ 1	Forward	5'-CTCCCGTGGCTTCTAGTGC-3'
	Reverse	5'-GCCTTAGTTTGGACAGGATCTG-3'
GAPDH	Forward	5'-CCAGGTTGTCTCCTGCGACTT-3'
	Reverse	5'-CCTGTTGCTGTAGCCGTATTCA-3'
Foxp3-promoter	Forward	5'-TTCCTCCCGCTCTCTGACTCT-3
	Reverse	5'-AAGCGCCAGTTGTGTACAAATATC-3'
Foxp3-CNS1	Forward	5'-GTTTTGTGTTTTAAGTCTTTTGCACCTTG-3'
	Reverse	5'-CAGTAAATGGAAAAAATGAAGCCATA-3'
<b>(For Taqman)</b>		<b>Cat#</b>
Hprt1		Mm00446968_m1
Foxp3		Mm00475162_m1
Tgfb1		Mm00441724_m1
Il10		Mm01288386_m1
Tnf		Mm00443258_m1
Tgfb1		Mm00436964_m1
Tgfb2		Mm03024091_m1
Smad6		Mm00484738_m1
Smad7		Mm00484742_m1
Rel		Mm01239661_m1
Tbx1		Mm00448949_m1
Rorc		Mm01261022_m1
Nfkb1		Mm00446968_m1
Nfkb2		Mm00476361_m1
Ikbkb		Mm01222247_m1
Jun		Mm00495062_s1
Il2ra		Mm01340213_m1
Il2rb		Mm00434268_m1
Smad2		Mm00487530_m1
Smad3		Mm00489637_m1
Smad4		Mm03023996_m1
Cd3e		Mm01179194_m1
Icos		Mm00497600_m1
Nfkbiz		Mm00600522_m1
Cd44		Mm01277161_m1
Stat5a		Mm03053818_s1
Stat5b		Mm0083989_m1
Jak1		Mm00600614_m1
Hdac4		Mm01299552_m1
Ifng		Mm01168134_m1
Il2		Mm00434256_m1
Il5		Mm00439646_m1
Il13		Mm00434204_m1
Inha		Mm00439683_m1
Rorc		Mm01261022_m1
Batf		Mm00479410_m1
Stat3		Mm00456961_m1

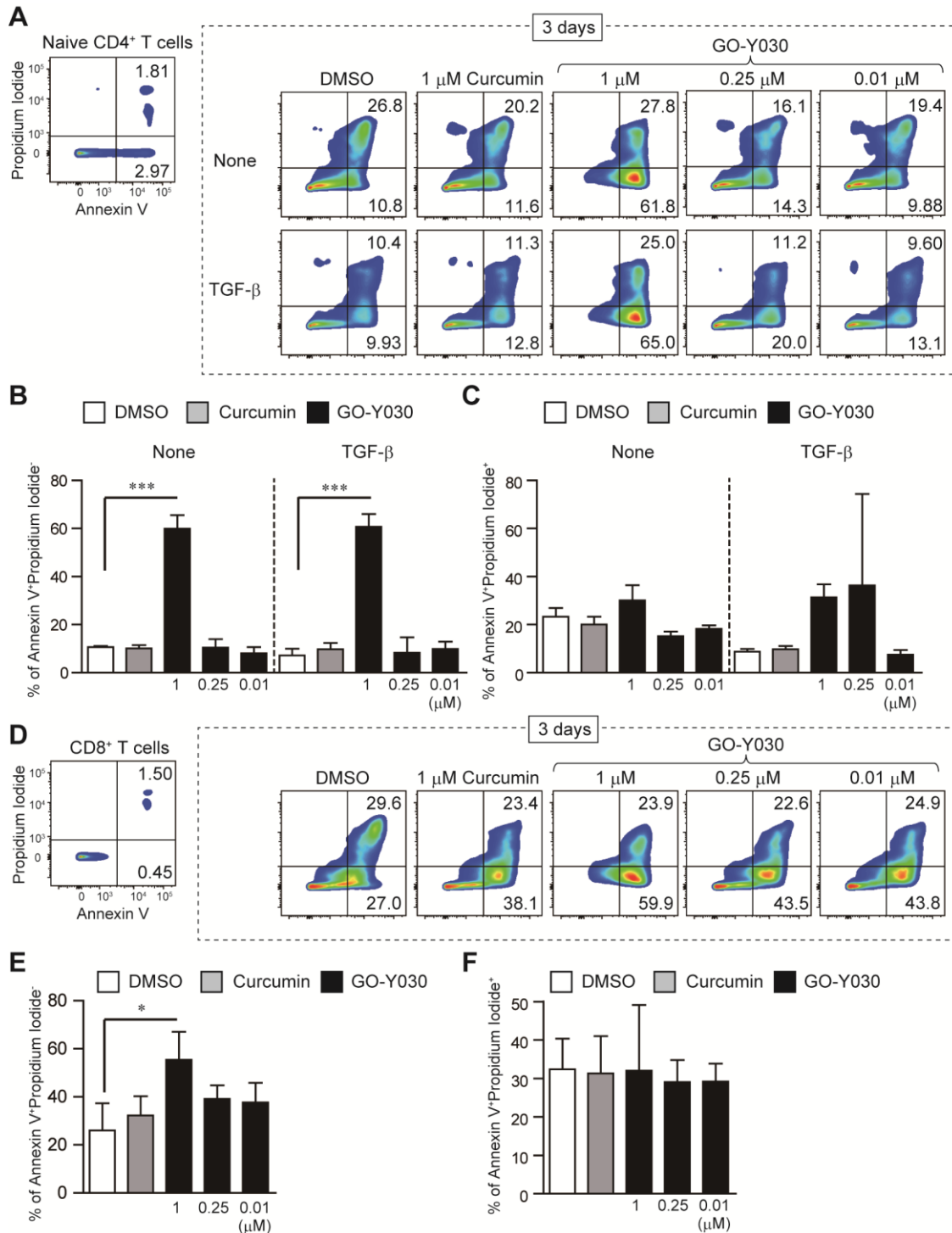
Myd88	Mm00440338_m1
Myc	Mm00487803_m1
Hk1	Mm00439344_m1
Hk2	Mm00443385_m1
Socs1	Mm00782550_s1
Socs3	Mm00545913_s1
Slc2a1	Mm00441480_m1
Slc2a3	Mm00441483_m1
Runx1	Mm01213404_m1
Runx3	Mm00490666_m1
Tcf12	Mm00441699_m1
Cd274	Mm03048248_m1
Rara	Mm01296312_m1
Rora	Mm01173766_m1

**Supplementary Table 1: Primers List**



**Supplementary Figure 1: Naive CD4<sup>+</sup> T cells from C57/BL6 mice cultured in each concentration of GO-Y-030.**

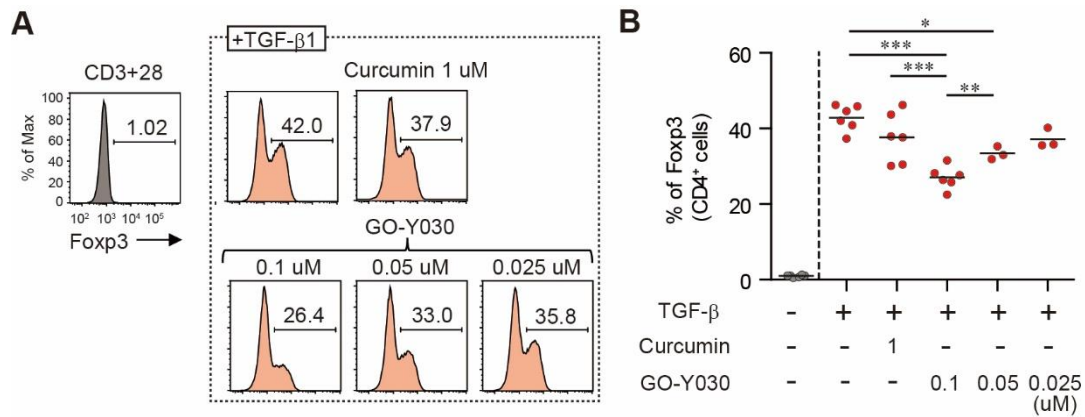
(A, B) Representative SSC and FSC FACS analysis at five independent experiments. Purified naïve CD4<sup>+</sup> T cells cultured for three days, and then gated live cell population according to SSC and FSC. Statistical analyses were performed in each concentration between Curcumin and GO-Y030, and 2 ng/mL TGF- $\beta$  versus all.



**Supplementary Figure 2: Effects of GO-Y-030 in T cell viability.**

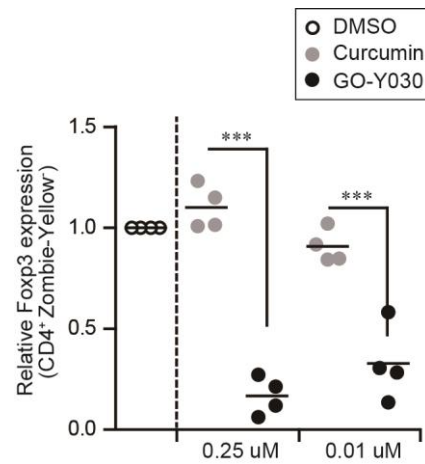
(A) Representative Annexin V and Propidium Iodide FACS analyses at three independent experiments. Purified naïve CD4<sup>+</sup> T cells cultured for three days, and then gated CD4<sup>+</sup> cell population. (B, C) Statistic analyses were performed in each concentration between Curcumin and GO-Y030 versus DMSO control. One-way analysis of variance (ANOVA) with post-hoc Tukey's multiple comparisons test employed. (D) Representative Annexin V and Propidium Iodide FACS

analyses at three independent experiments. Purified naïve CD8<sup>+</sup> T cells cultured for three days, and then gated CD8<sup>+</sup> cell population. **(E, F)** Statistic analyses were performed in each concentration between Curcumin and GO-Y030 versus DMSO control. One-way ANOVA with post-hoc Tukey's multiple comparisons test employed.



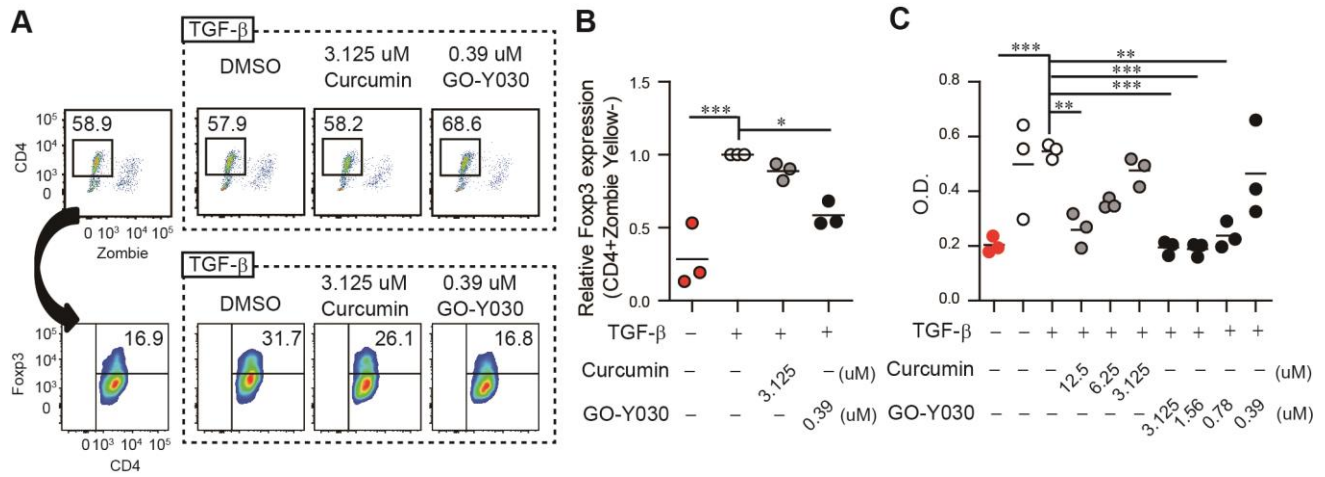
### Supplementary Figure 3: GO-Y030 prevent TGF- $\beta$ induced Foxp3<sup>+</sup>Tregs generation

(A, B) Frequency of Foxp3<sup>+</sup> Tregs in total CD4<sup>+</sup> cells. Splenic naïve CD4<sup>+</sup>T cells were cultured in the presence or absence of 2 ng/mL TGF- $\beta$ , 1  $\mu$ M curcumin, or 0.1-0.025  $\mu$ M GO-Y030 for three days. One-way analysis of variance (ANOVA) with post-hoc Tukey's multiple comparisons test employed.



**Supplementary Figure 4: Relative Foxp3<sup>+</sup>Tregs in CD4<sup>+</sup>Zombie Yellow<sup>-</sup> T cells.**

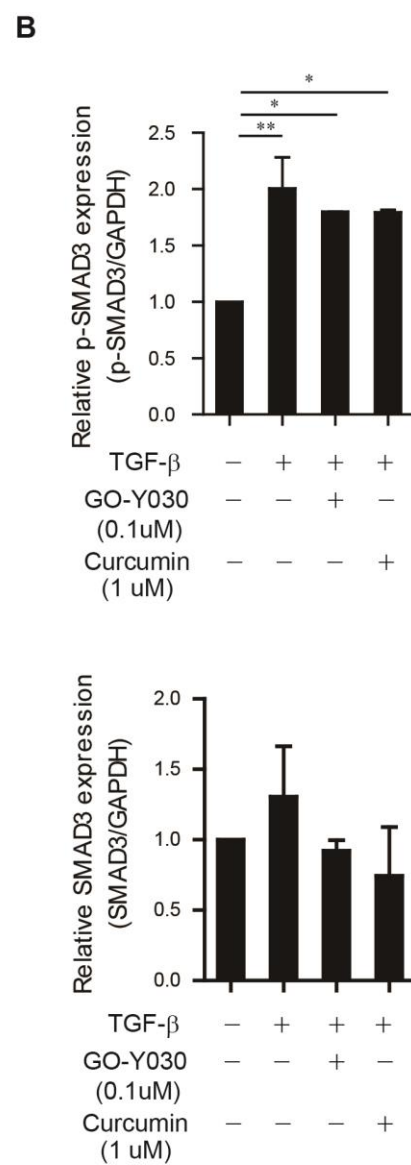
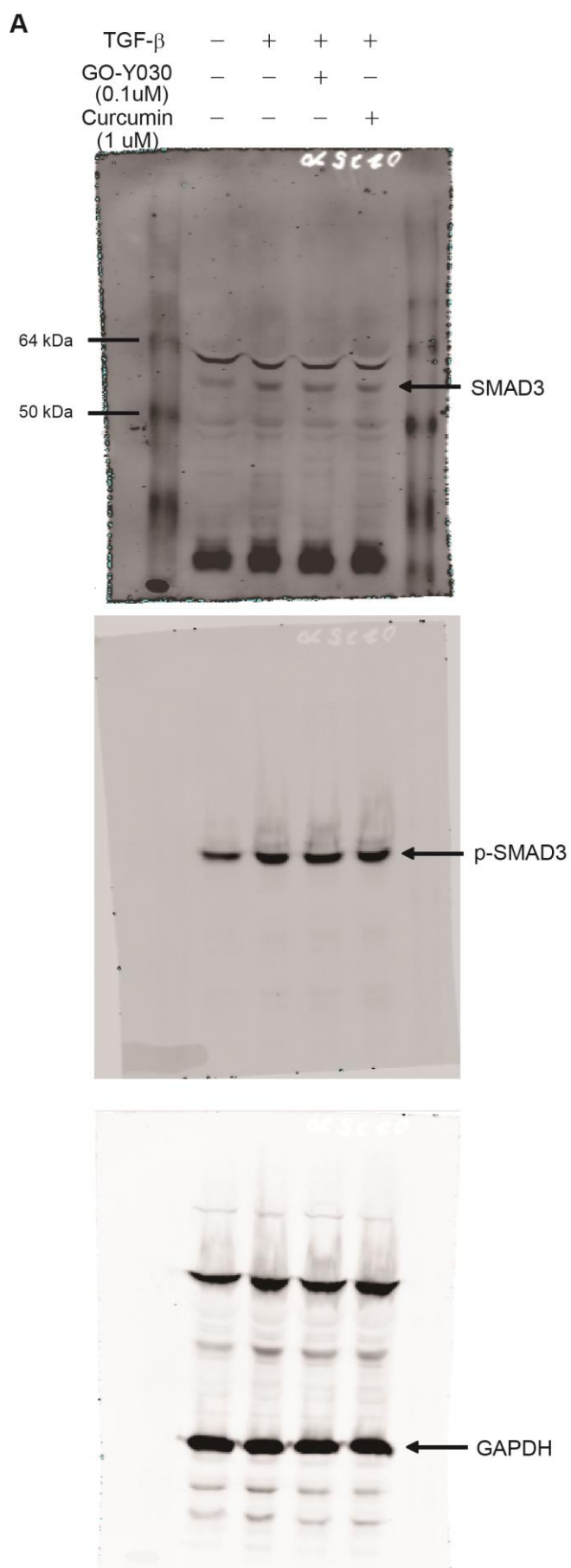
Splenic naïve CD4<sup>+</sup> T cells were cultured in the presence of 2 ng/mL TGF- $\beta$  with or without curcumin or GO-Y030 for three days. Percentage of TGF- $\beta$ -induced Foxp3<sup>+</sup> population is as set as “1”. One-way analysis of variance (ANOVA) with post-hoc Tukey’s multiple comparisons test employed.



**Supplementary Figure 5: GO-Y030 prevents TGF- $\beta$ -induced Foxp3<sup>+</sup>Tregs in human naïve CD4<sup>+</sup> T cells.**

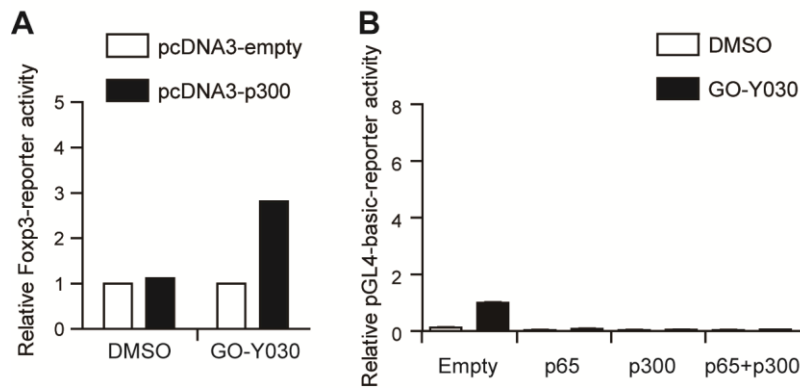
(A) Human naïve CD4<sup>+</sup>T cells were cultured in the presence of 2 ng/mL TGF- $\beta$ 1 with or without curcumin or GO-Y030 for three days. Data are one representative at three independent experiments. (B) Relative Foxp3 expression in CD4<sup>+</sup> T cells. TGF- $\beta$ 1 stimulation only is set as “1”. One-way analysis of variance (ANOVA) with post-hoc Dunnet’s multiple (vs. TGF- $\beta$ ) comparisons test employed. (C) Relative live cell counts. Human naïve CD4<sup>+</sup> T cells were cultured with or without 2 ng/mL TGF $\beta$  and concentrations of Curcumin or GO-Y030 as indicated for 72 h followed by the addition of the cell counting reagent. Red: No cells (Medium alone). Data are representative at three independent experiments using different healthy donors. One-way ANOVA with post-hoc Dunnet’s multiple (vs. TGF- $\beta$ ) comparisons test employed.





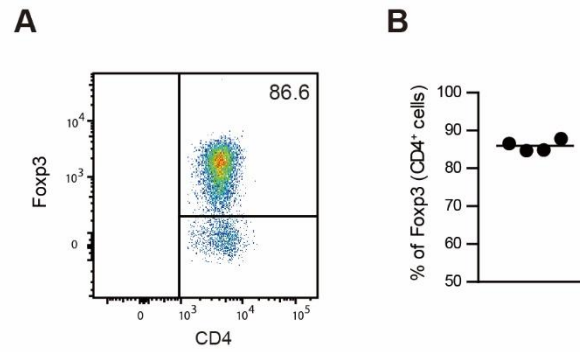
**Supplemental Figure 6: GO-Y030 does not affect TGF- $\beta$ -induced SMAD pathway.**

(A) Representative western blotting image of SMAD3, phospho-SMAD3 and GAPDH at three independent experiments. Naive CD4<sup>+</sup> T cells were stimulated with or without TGF- $\beta$  in the presence or absence of 0.1  $\mu$ M GO-Y030 or 1  $\mu$ M Curcumin. (B) Relative SMAD3 and phospho-SMAD3 expression. (n=3, Mean with standard error of the mean) Without TGF- $\beta$  and GO-Y030 stimulation was set as “1”. One-way analysis of variance (ANOVA) with post-hoc Tukey’s multiple comparisons test employed. One-way ANOVA with post-hoc Tukey’s multiple comparisons test employed.



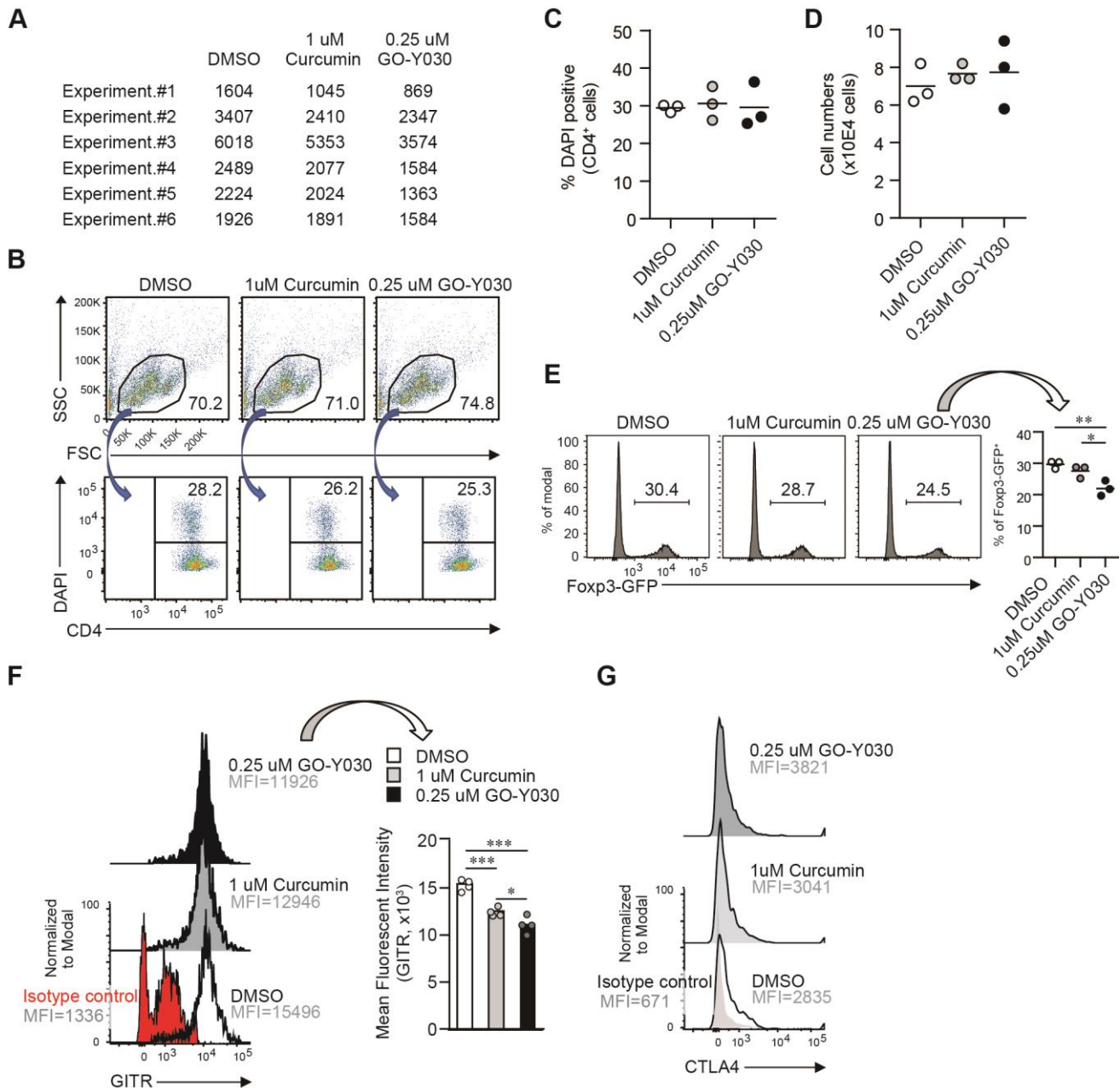
### Supplemental Figure 7: Relative murine Foxp3-promoter activity

(A) pGL4-Foxp3 promoter (-1702 to +174) activity were analyzed using the Duo-luciferase assay systems. DMSO- or 1.0  $\mu$ M GO-Y030 were treated HEK293 before 24 h electroporation. pcDNA3-empty vector transfection (Firefly/Renilla) was set as “1”. (B) pGL4-basic promoter (control) activity was analyzed using the Duoluciferase assay systems. DMSO- or 1.0  $\mu$ M GO-Y030 were treated HEK293 cells before 24 h transfection. Data are one representative at two independent experiments (n=2, Mean + standard deviation).



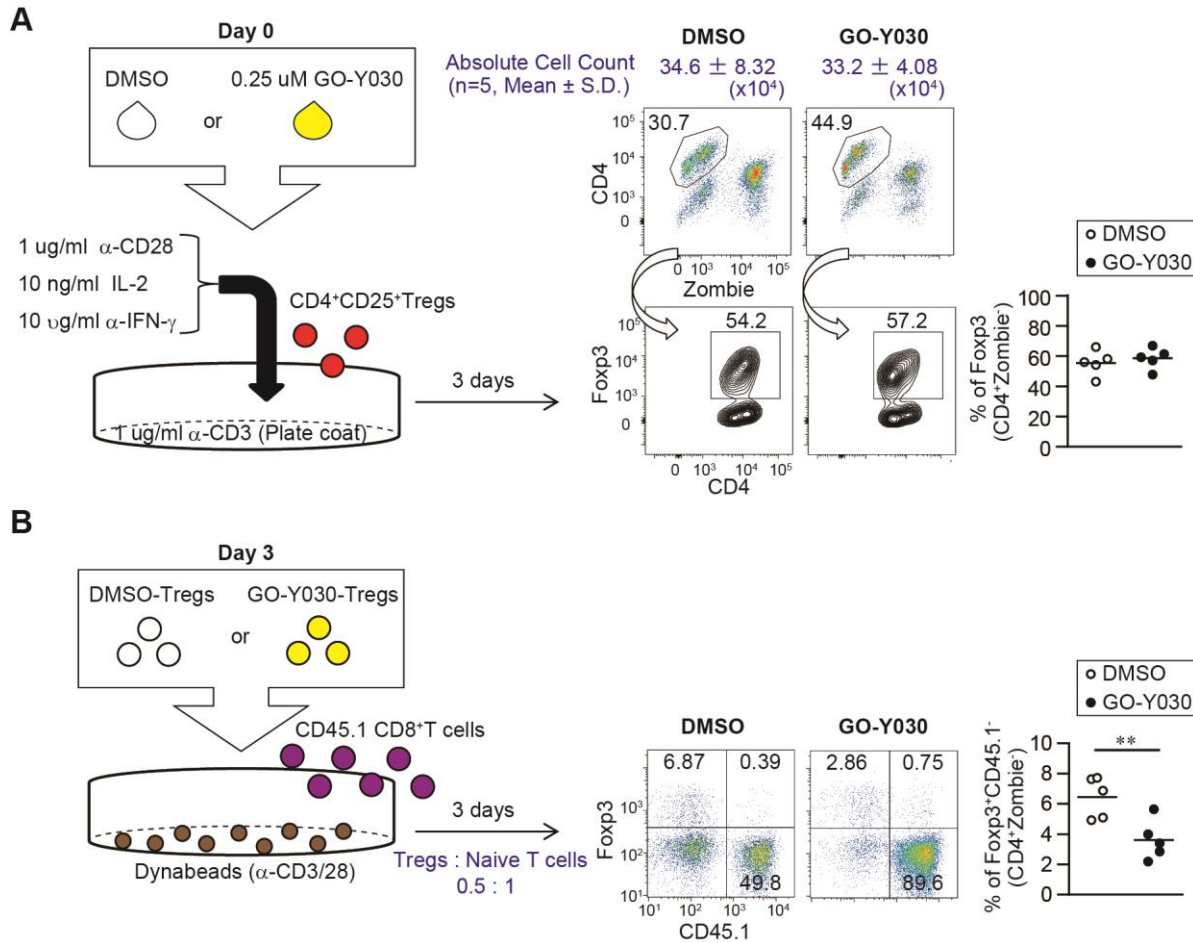
**Supplemental Figure 8: Representative Foxp3 expression in Tregs after CD4<sup>+</sup>CD25<sup>+</sup> isolation.**

(A, B) Splenic CD4<sup>+</sup>CD25<sup>+</sup> Tregs were isolated by using autoMACS (Milteny Biotech) and FACS analyses. Data are one representative of three independent experiments.



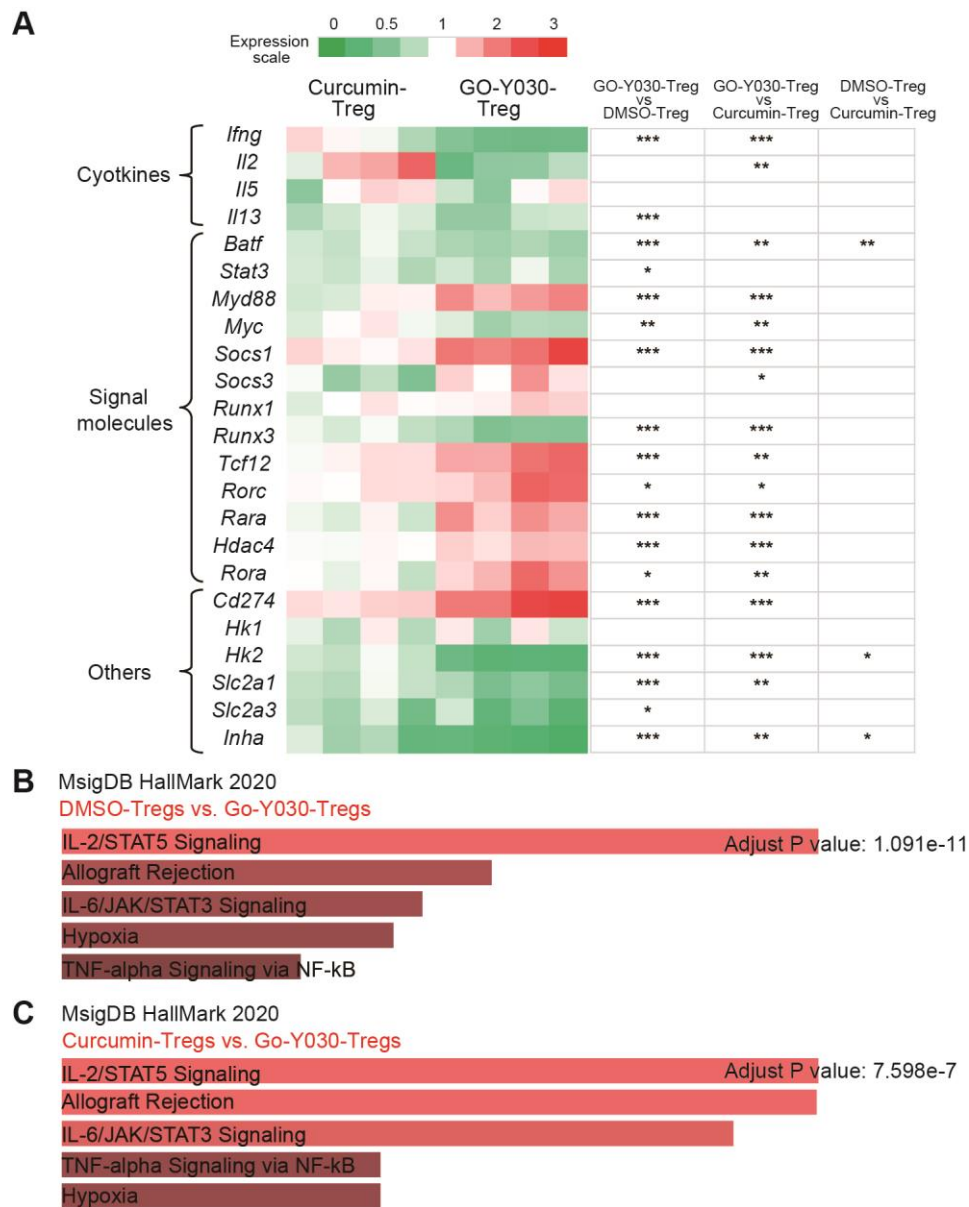
### Supplemental Figure 9: Phenotype of Cultured CD4<sup>+</sup>CD25<sup>+</sup> Tregs.

(A) Foxp3 Mean fluorescence Intensity in CD4<sup>+</sup>Foxp3<sup>+</sup> Treg populations in Figure.3A. Data shows six independent experiments. experiments. (B, C) Relative cell survival rate after 18 h culture of CD4<sup>+</sup>CD25<sup>+</sup> Tregs with or without Curcumin or GO-Y030. Data are one representative at more than three independent experiments. (D) Absolute number of cells after 18 h culture of CD4<sup>+</sup>CD25<sup>+</sup> Tregs with or without Curcumin or GO-Y030. The starting number of cells in each wells was 1 x 10<sup>5</sup> cells. (E) Foxp3-GFP positive cells in CD4<sup>+</sup> population. Foxp3-GFP positive cells were purified by FACS-Aria II (>95%) and cultured 18 h with CD3 + CD28. Data are one representative at three independent experiments. GITR (F) and CTLA4 (G) in CD4<sup>+</sup>Foxp3<sup>+</sup> Treg populations in Figure. 5A. Gray; Isotype control, Black; GITR or CTLA4. Data are one representative at three independent experiments. One-way analysis of variance with Tukey employed for statistic difference.



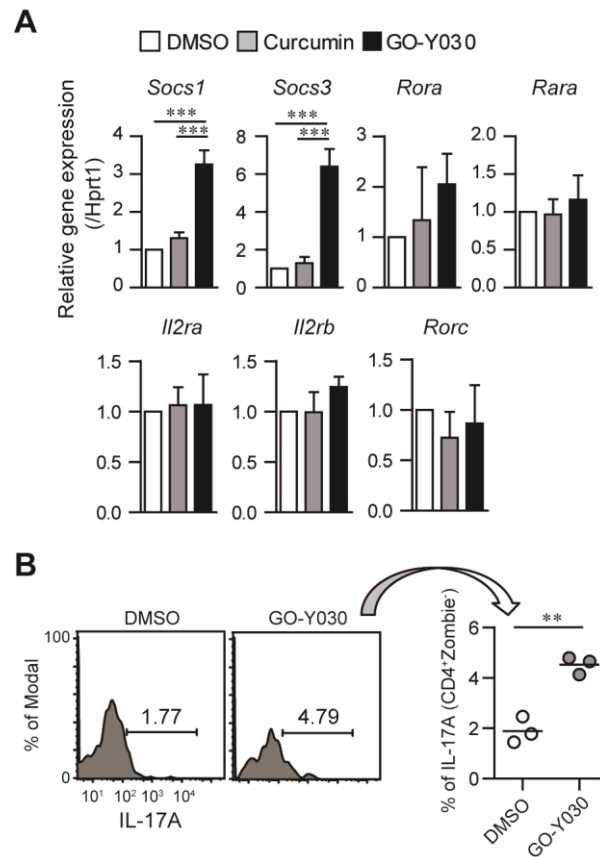
**Supplemental Figure 10: GO-Y030 controls IL-2/STAT5 axis in CD4<sup>+</sup>CD25<sup>+</sup> Tregs.**

(A) Percentage of Foxp3 in cultured CD4<sup>+</sup>CD25<sup>+</sup>Tregs at day three. Data are representative at five independent experiments. Student T-test was employed. (B) Percentage of Foxp3 in cultured CD4<sup>+</sup>CD25<sup>+</sup> Tregs in the suppression assay. DMSO- or GO-Y030-treated CD4<sup>+</sup>CD25<sup>+</sup> Tregs were co-cultured with CD8<sup>+</sup> T cells (Tregs:CD8<sup>+</sup> T cells=0.5:1) for three days. Data are representative at shows five independent experiments. Student T-test was employed.



**Supplemental Figure 11: GO-Y030 controls IL-2/STAT5 axis in CD4<sup>+</sup>CD25<sup>+</sup> Tregs.**

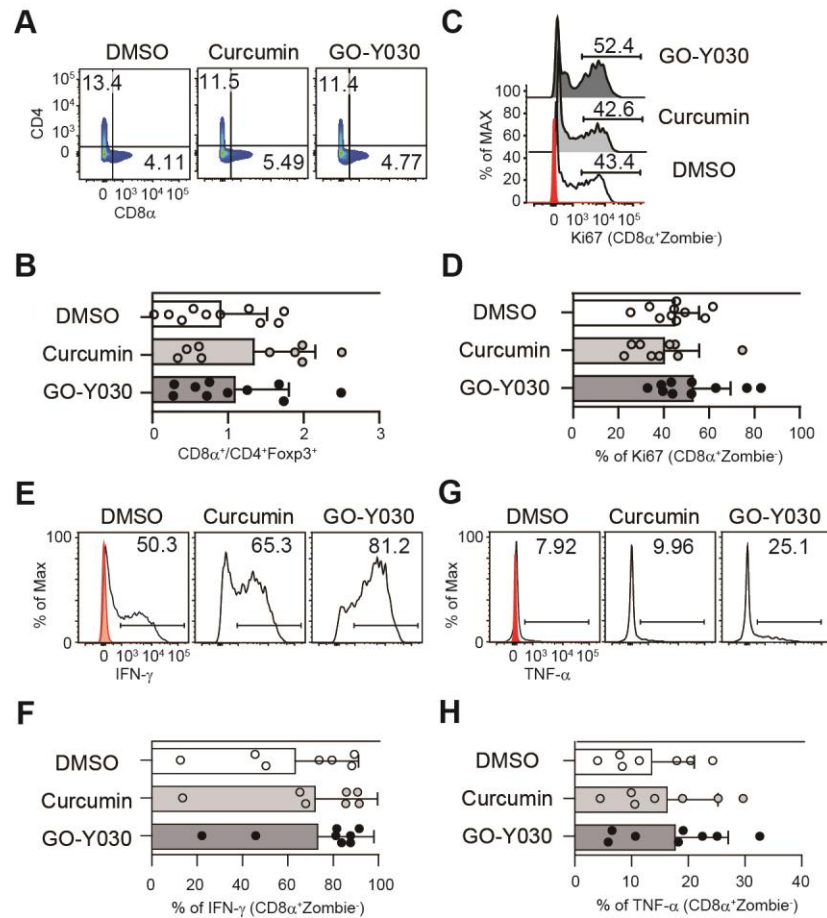
(A) Real time PCRs in 72 h culture of CD4<sup>+</sup>CD25<sup>+</sup> Tregs with or without Curcumin or GO-Y030. The color scale is shown at the top of heat map. Each genes expression of DMSO-treated Tregs are as set as “1”. Data showed four independent experiments. (B, C) Enrichr- was used to calculate enrichment scores of signaling pathways. We selected genes of significantly difference expression ( $P < 0.05$ ) between DMSO-Tregs and GO-Y030-Tregs (B, <https://maayanlab.cloud/Enrichr/enrich?dataset=ce1ae4783a07360aa829bddb0fd36eb1>) or Curcumin-Tregs and GO-Y030-Tregs (C, <https://maayanlab.cloud/Enrichr/enrich?dataset=35a8b176b017fbd7c09576aa75995cd>). Statistical analyses (One-way analysis of variance with Tukey) were performed.



**Supplemental Figure 12: GO-Y030 controls IL-2/STAT5 axis in CD4<sup>+</sup>Foxp3-GFP<sup>+</sup> Tregs.**

(A) Real time PCRs in 72 h culture of CD4<sup>+</sup>Foxp3-GFP<sup>+</sup> Tregs with or without 1  $\mu$ m Curcumin or 0.25  $\mu$ m GO-Y030. Data pooled three independent experiments. Statistical analyses (One-way analysis of variance with Tukey) were performed. (B). Th17 population in cultured CD4<sup>+</sup>Foxp3-GFP<sup>+</sup> Tregs with or without 0.25  $\mu$ m GO-Y030. Student T-test was performed.





**Supplemental Figure 13: GO-Y030 did not prevent infiltration and activation of CD8<sup>+</sup> cells in tumor microenvironment.**

(A) Frequency of CD4<sup>+</sup>/CD8α<sup>+</sup> cells in tumor infiltrate lymphocytes. (B) Ratio of tumor infiltrate CD8<sup>+</sup> cells to CD4<sup>+</sup>Foxp3<sup>+</sup> Treg cells. (C, D) Ki67 expression in CD8α<sup>+</sup> cells in tumor infiltrate lymphocytes. Red; isotype control. Data are one representative of each of the two independent experiments (A, C). (E, F) IFN-γ production from CD8α<sup>+</sup> cells in tumor infiltrate lymphocytes. Red; isotype control. (G, H) TNF-α production from CD8α<sup>+</sup> cells in tumor infiltrate lymphocytes. Red; isotype control. One-way analysis of variance with post-hoc Tukey's multiple comparisons test was used (B, D, F, H). The graph shows mean and standard deviation.