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PPARγ Negatively Regulates T Cell Activation to Prevent Follicular Helper T Cells and Germinal Center Formation



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Abstract

Peroxisome proliferator-activated receptor gamma (PPAR γ) is a transcription factor that regulates lipid and glucose metabolism. Although studies of PPAR γ ligands have demonstrated its regulatory functions in inflammation and adaptive immunity, its intrinsic role in T cells and autoimmunity has yet to be fully elucidated. Here we used CD4-PPAR γ^{KO} mice to investigate PPAR γ -deficient T cells, which were hyper-reactive to produce higher levels of cytokines and exhibited greater proliferation than wild type T cells with increased ERK and AKT phosphorylation. Diminished expression of IkB α , Sirt1, and Foxo1, which are inhibitors of NF- κ B, was observed in PPAR γ -deficient T cells that were prone to produce all the signature cytokines under Th1, Th2, Th17, and Th9 skewing condition. Interestingly, 1-year-old CD4-PPAR γ^{KO} mice spontaneously developed moderate autoimmune phenotype by increased activated T cells, follicular helper T cells (T_{FH} cells) and germinal centers in CD4-PPAR γ^{KO} mice and the T cells showed increased of Bcl-6 and IL-21 expression suggesting its regulatory role in germinal center reaction. Collectively, these results suggest that PPAR γ has a regulatory role for T_{FH} cells and germinal center reaction to prevent autoimmunity.

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Introduction

Nuclear receptors constitute a superfamily of ligand-dependent transcription factors that regulate diverse aspects of metabolism and homeostasis. This family is further subdivided into three subclasses based on their DNA-binding and ligand-binding properties, including classical steroid hormone receptors, adopted orphan receptors, and orphan receptors [1-4]. Peroxisome Proliferator-activated Receptors (PPAR) are emerging adopted orphan nuclear receptors that regulate lipid and glucose metabolism, as well as cell growth, differentiation, apoptosis, and immunity, in a wide variety of cells. The PPAR subfamily consists of three subtypes including PPAR α , PPAR β/δ , and PPAR γ . Each member binds multiple ligands, regulates target genes, and plays different biological roles. The ligands for PPAR γ have been used for therapeutic applications to treat atherosclerosis, obesityinduced insulin resistance, and inflammatory bowel disease [5-7]. For example, synthetic PPAR γ agonists called thiazolidinediones (TZD) improved insulin sensitivity in metabolic syndrome [5,8]. The PPAR γ agonist 15-deoxy- $\Delta^{12,14}$ -prostaglandin J₂ (15-PGJ₂) has been shown to downregulate T cell proliferation and IL- 2 production [9]. Another PPAR γ agonist, ciglitazone, has been shown to increase regulatory T (Treg) cell differentiation from naïve T cells [10], while pioglitazone has been demonstrated to ameliorate EAE severity through selective inhibition of Th17 differentiation [11] and inhibit human alloresponses in a humanized model of graft arteriosclerosis [12]. More recently, PPAR γ has been shown to regulate Treg accumulation in visceral adipose tissue, and pioglitazone treatment was required for complete recovery from insulin sensitivity in obese mice [13].

Autoimmune disease occurs when the immune system recognizes self-molecules and responds by inducing autoantibody production, germinal center (GC) formation, and chronic inflammation [14–16]. Follicular helper T (T_{FH}) cells are involved in the antibody response by providing selection signals to GC B cells and promoting plasma and memory cell differentiation [17]. Bcl-6 mediates the development of T_{FH} cells, which are characterized by expression of the chemokine receptor CXCR5, ICOS, and PD-1 [18–21]. Recent studies have suggested T_{FH} cells as a potential therapeutic target for treating autoimmune diseases [15,22,23].

Although previous studies focused on PPAR γ agonists revealed its regulatory function in T cells, the mechanism and intrinsic roles of this nuclear receptor in T_{FH} cells and autoimmunity is largely uncharacterized. Therefore, we investigated the role of PPAR γ in T cells using CD4 T cell-specific PPAR γ knockout mice. Interestingly, we found that PPAR γ -deficient T cells exhibit enhanced proliferation following T cell receptor (TCR) stimulation and increased cytokine production under Th1, Th2, Th17, and Th9 differentiation conditions. Interestingly moderate autoimmune phenotypes developed spontaneously in 1-year-old CD4-PPAR γ^{KO} mice with increased activated T cells, T_{FH} cells, and GCs. Furthermore, PPAR γ regulated T cell activation by modulating the stability of IkB α , Foxo1, and Sirt1, which are negative regulators of NF- κ B and T_{FH} cells. These results suggest the important regulatory roles of PPAR γ in T_{FH} cells and GC formation in the prevention of autoimmunity.

Results

$\ensuremath{\text{PPAR}\gamma}$ is highly expressed in CD4 T cells upon TCR stimulation

To determine its expression in various lymphocyte subsets, the level of PPAR γ mRNA was examined in FACS-purified CD4⁺ T, CD8⁺ T, and CD19⁺ B cells by quantitative real-time PCR analysis. PPAR γ expression was significantly higher in CD4⁺ T cells compared to CD8⁺ T cells and CD19⁺ B cells (Figure 1A), suggesting that this nuclear receptor may play important roles in CD4⁺ T cells. In addition, PPAR γ mRNA expression was significantly increased following TCR stimulation (Figure 1B), further implying its regulatory role in controlling activated CD4⁺ T cell functions.

Next, we generated mice in which PPAR γ was specifically deleted in CD4⁺ T cells by breeding PPAR $\gamma^{fl/fl}$ with CD4-Cre transgenic mice (Figure 1C&D). PPAR γ -deficiency did not affect the frequency of CD4⁺, CD8⁺, and Foxp3⁺ natural regulatory T cell (nTreg) in the thymus of 6- to 8-week-old mice (Figure S1A-D), the proportion of lymphocyte population including B cells, NK cells, and naïve or memory T cells in the spleen (Figure S1E-L). Altogether these data suggest that PPAR γ deletion in CD4⁺ T cells does not affect thymic and splenic cell populations that PPAR γ is dispensable for T cell development.

$PPAR\gamma$ -deficient T cells are hyper-reactive to TCR stimulation

Although PPAR γ is dispensable for lymphocyte development, increased PPAR γ expression following TCR stimulation led us to examine its function during T cell activation. To address this, we examined cytokine production from purified CD4⁺CD25⁻ T cells stimulated with anti-CD3 and anti-CD28 antibodies. Various cytokines including IFN- γ , IL-4, IL-17, and IL-2 were increased in PPAR γ -deficient T cells compared to littermate control T cells, (Figure 2A-D) suggesting that these cells are hyper-reactive to TCR stimulation. In addition, PPAR γ -deficient T cells proliferated significantly more than control CD4⁺ T cells (Figure 2E), suggesting that PPAR γ plays a role as a negative regulator of T cell activation and proliferation.

Next, we investigated cytokine production after CD4⁺CD25⁻ T cells were differentiated into various helper T cell subsets including Th1, Th2, Th17, and Th9. Interestingly, production of all the signature cytokines secreted by each helper T cell subset, namely IFN- γ , IL-13, IL-17, and IL-9, respectively, were significantly increased in PPAR γ -deficient T cells (Figure 2F) without affecting differentiation efficiency (Figure 2G). These results suggest that PPAR γ has a negative regulatory role for T cell activation and proliferation.

PPAR γ maintains the stability of I κ B α , Foxo1, and Sirt1 to regulate TCR signaling

A previous study reported that PPAR γ inhibits NF- κ B in macrophages [24]. However, the molecular basis of how PPAR γ regulates T cell activation has not been fully elucidated. To determine the mechanism, we assessed ERK and AKT phos- $CD4^+$ phorylation in splenic naïve Т cells (CD4⁺CD25⁻CD62L^{high}) stimulated with anti-CD3/CD28-coated beads. Following TCR stimulation for 20 min, the PPARydeficient CD4+ T cells showed increased ERK and AKT phosphorylation compared to littermate control T cells (Figure 3A), suggesting that $PPAR\gamma$ inhibits events of TCR downstream signaling pathway.

We performed microarray analysis to identify possible target genes (e.g., Foxo1) regulated by PPAR γ (data not shown). Interestingly, TCR stimulation of PPAR γ -deficient CD4⁺ T cells resulted in reduced expression of I κ B α , Foxo1, and Sirt1 (Figure 3B), which have been recently identified as negative regulators of T cell activation [25,26]. In addition, phosphorylated p65 level was increased in PPAR γ -deficient CD4⁺ T cells suggesting that PPAR γ contributes to the stability of I κ B α , Foxo1, and Sirt1 to regulate NF- κ B in activated T cells.

PPAR γ deficiency in T cells spontaneously enhances activated T cells in vivo

The TCR-stimulated hyper-reactivity observed in PPARydeficient T cells from 6- to 8-week-old CD4-PPARyKO mice prompted us to investigate spontaneous changes in the T cell phenotype of 1-year-old CD4-PPAR γ^{KO} mice. CD62L is a cell adhesion molecule, which is highly expressed on naïve T lymphocytes. The frequency of CD4⁺CD62L^{low} cells was increased significantly in the mesenteric lymph nodes (MLN) of CD4-PPAR γ^{KO} mice compared to age-matched control mice (Figure 4A&B). The proportion of CD4⁺CD62L^{low} cells of CD4-PPARy^{KO} mice was augmented compared to littermate control. These results are supported by an increase of the early activation marker CD69 expressing CD4⁺ T cells (Figure 4C&D). Additionally, we investigated the expression of IL-7R α , which has been shown to be decreased in activated T cells [27]. PPAR γ deficiency in T cells from 1-year-old CD4-PPAR γ^{KO} mice showed significantly reduced proportion of CD4⁺IL-7R α^{high} cells in the MLN (Figure 4E&F). Collectively, these data demonstrate that $PPAR\gamma$ deficiency in T cells contributes to their hyper-reactivity, which induces spontaneous T cell activation in vivo.

PPARγ-deficient T cells in mice develop a spontaneous autoimmune phenotype

Because the number of activated T cells was increased in 1-yearold CD4-PPAR γ^{KO} mice, we next investigated autoimmune phenotype in these animals. Flow cytometric analysis of splenocytes from 6-month-old CD4-PPAR γ^{KO} mice demonstrated that the proportion of CD19⁺ B cells was increased while CD4⁺ T cells were slightly decreased compared to littermate control mice (Figure 5A). Based on this observation, we hypothesized that the hyper-reactive PPAR γ -deficient T cells promote greater B cell activation, thereby leading to autoantibody production. Our data reveal that the proportion of CD19⁺ and CD138⁺ plasma cells was increased in the spleen of CD4-PPAR γ^{KO} mice (Figure 5B&C). Moreover, the level of anti-nuclear antibodies (ANA) in serum from 6-month-old mice (Figure 5D) and the level of anti-doublestranded DNA (dsDNA) antibodies in serum from 1-year-old mice (Figure 5E) were higher than age-matched control mice.



Figure 1. PPAR γ **expression in T cells and generation of CD4-PPAR** γ^{KO} **mice.** (A) Expression of PPAR γ in FACS-purified CD4⁺, CD8⁺, and CD19⁺ cells from female wild type C57BL/6J mice was analyzed by quantitative real-time PCR and normalized to β -actin. Values represent the mean \pm SEM, n = 4. **P<0.01 when CD19 and CD8 were compared. (B) MACS-purified naïve (CD4⁺CD25⁻CD62L^{high}CD44^{low}) T cells from female wild type C57BL/6J mice were stimulated with plate-bound anti-CD3 and soluble anti-CD28 antibodies for the indicated times. PPAR γ expression was determined by quantitative real-time PCR. Relative PPAR γ expression level was calculated as the fold-change relative to 0 h. Values represent the mean \pm SEM, n = 4. *P<0.05, **P<0.01 when 0 h, 24 h, and 48 h were compared. (C) FACS-sorted cells from female littermate control (Cre-) and CD4-PPAR γ^{KO} (Cre+) mice were harvested and PPAR γ expression was determined by quantitative real-time PCR. Values represent the mean \pm SEM, n = 4. *P<0.05, **P<0.01 when 0 h, 24 h, and 48 h were compared. (C) FACS-sorted cells from female littermate control (Cre-) and CD4-PPAR γ^{KO} (Cre+) mice were harvested and PPAR γ expression was determined by quantitative real-time PCR. Values represent the mean \pm SEM, n = 3. *P<0.05. (D) Cre-mediated PPAR γ deletion in cell lysate prepared from CD4⁺ T cells was confirmed by western blot analysis. doi:10.1371/journal.pone.0099127.g001

To further examine the autoimmune phenotype, inflammation in the kidneys was examined and scored by histologic analysis of hematoxylin and eosin-stained sections (Figure 5F&G). Kidney from 1-year-old CD4-PPAR γ^{KO} mice exhibited an expanded double layer around the glomerulus with thicker tubules compared to age-matched control mice. These results suggest that PPAR γ deficiency in T cells leads to autoantibody production and spontaneous autoimmune disease in aged mice.

$\mathsf{PPAR}\gamma\text{-deficient}\ \mathsf{T}\ \mathsf{cells}\ \mathsf{contribute}\ \mathsf{to}\ \mathsf{T}_{\mathsf{FH}}\ \mathsf{cells}\ \mathsf{and}\ \mathsf{GC}$ development

 $T_{\rm FH}$ cells are crucial for providing help to B cells within the germinal center, where somatic hyper-mutation, class-switch recombination, and differentiation of memory B cells and long-lived plasma cells occur [28–31]. Due to increased B cell numbers, including plasma cells, in 6-month-old CD4-PPAR $\gamma^{\rm KO}$ mice, we hypothesized that PPAR γ deficiency in T cells lead to increased T_{\rm FH} cells *in vivo*. Previous report has shown that PSGL-1 low population enriched more $T_{\rm FH}$ cells than PSGL-1 high [31] so that PSGL-1^{low} gating strategy was applied to analyze T_{\rm FH} cells.

Flow cytometric analysis revealed an increased proportion of CXCR5⁺PD-1⁺ T_{FH} cells in 1-year-old CD4-PPAR γ^{KO} mice (Figure 6A - C). The number of GCs found in the spleen of these animals was also increased compared to age-matched control mice (Figure 6D&E). In addition, proportion of germinal center B cells (CD95⁺GL7⁺ from B220⁺ cells) were increased in the spleen with elevated amount of IgG1 and reduced IgM from the sera in 1-year-old CD4-PPAR γ^{KO} mice (Figure 6F&G). These results suggest that the spontaneous autoimmune phenotypes observed were related to increased T_{FH} cells and enhanced GC reaction.

To further confirm that PPAR γ regulates induction of T_{FH} cells and GC reaction, 8-week-old mice were immunized with sheep red blood cells (SRBC) and the T_{FH} cells and GC number were analyzed on day 7 (Figure 7A). Following SRBC immunization, the proportion of CXCR5⁺PD-1⁺ T_{FH} cells (Figure 7B&C) and total number of spleen cells were significantly increased (Figure 7D) which is correlated to enhanced GC number (Figure 7E&F) in CD4-PPAR γ^{KO} mice than in control animals. Next, we examined IL-21 and Bcl-6 expression levels, which are important molecules for induction of T_{FH} cells and GC reaction. CD4-PPAR γ^{KO} T



Figure 2. PPAR γ **deficiency induces hyper-reactivity in T cells.** CD4⁺CD25⁻ T cells from the spleens of 6- to 8-week-old female littermate control (Cre⁻) and CD4-PPAR γ^{KO} (Cre+) mice were stimulated for 24 h with plate-bound anti-CD3 and soluble anti-CD28 antibodies. (A) IFN- γ , (B) IL-4, (C) IL-17, and (D) IL-2 levels in culture supernatants were measured by ELISA. (E) Proliferation of 3-day-anti-CD3/CD28 stimulated cells was measured by H³-thymidine incorporation. Values represent the mean ± SEM of counts per minute (CPM) in triplicate wells. CD4⁺25⁻ T cells were differentiated under Th1, Th2, Th17, and Th9 differentiation conditions. (F) After 5 days, the culture supernatants were collected and cytokine levels (IFN- γ , IL-13, IL-17, and IL-9) were measured by ELISA. Values represent the mean ± SEM, n = 4–6. **P*<0.05. (G) Flow cytometric analysis for lineage-specific cytokines (IFN- γ , IL-17, and IL-9) of Th1, Th2, Th17 and Th9 were determined by intracellular staining. Representative data were shown from five independent experiments. doi:10.1371/journal.pone.0099127.g002

cells expressed significantly higher level of IL-21 and Bcl-6 (Figure 7G&H) while they have reduced level of Blimp-1 (Figure 7I) in Th2 or Th17 cells suggesting PPAR γ negatively regulates Bcl-6 and IL-21 to inhibit T_{FH} cell differentiation and GC reaction. These results suggest that PPAR γ inhibits formation

of $T_{\rm FH}$ cells and GC reaction via regulation of Bcl-6 and IL-21 to prevent autoimmune disease.

Discussion

Based on its ligand-dependent functions, PPAR γ has been demonstrated to be an important negative regulator in T cells,

A		С	re-		Cre+				
	0	5	10	20	0	5	10	20	(min)
	an and			-		-			pERK
	1.00	1.46	2.68	5.95	1.29	1.22	4.57	6.06	
	4.00	0.40	4.00	0.54	0.00	4.00	7.40	40.00	pAKT
	1.00	2.48	4.08	9.54	2.68	4.90	7.10	12.62	Actin
В		0				C	ro		
			10-						(
	0	5	10	20	0	5	10	20	(min)
				-	-			-	p-p65
	1.00	0.59	1.10	1.21	1.35	1.12	1.72	2.68	l. n
	1 00	1 10	1 15	1 21	1 17	1 17	0.01	0.62	ΙκΒα
			-		-	-		0.02	Foxo1
	1.00	1.08	0.96	1.02	0.83	0.65	0.54	0.19	_
			-	-		-			Sirt1
	1.00	0.91	0.75	0.82	0.62	0.47	0.45	0.47	Actin

.

Figure 3. PPARγ **deficiency in T cells reduces the expression of molecules that inhibit NF-κB.** MACS-purified CD4⁺CD25⁻CD62L^{high} naive T cells from the spleens of 6- to 8-week-old female littermate control (Cre–) and CD4-PPARγ^{KO} (Cre+) mice were stimulated with anti-CD3 and anti-CD28 Dynabeads for the indicated times. (A) Levels of ERK and AKT phosphorylation and (B) Levels of p65 phosphorylation, IkBα, Foxo1, and Sirt1 were assessed by western blot analysis. The arbitrary unit represents band intensity normalized to β-actin and the relative value to WT 0 min is indicated. Representative data were shown from at least two independent experiments. doi:10.1371/journal.pone.0099127.q003

macrophages, and dendritic cells [9,24]. Previous studies have also suggested that PPAR γ is an important therapeutic target for treating autoimmune disease [5–7]. Here, for the first time, we characterized CD4-PPAR γ^{KO} mice to investigate the intrinsic role of PPAR γ in T cells and autoimmunity related to T_{FH} cells and GC reaction.

We found that, among lymphocytes, PPAR γ was expressed predominantly in CD4⁺ T cells. The expression level of PPAR γ , however, increased following TCR stimulation. This is consistent with a previous study that reported an increase in PPAR γ expression following anti-CD3 antibody stimulation and PPAR γ ligand (15d-PGJ2 and ciglitazone) treatment, which it also contributed to the inhibition of T cell proliferation and IL-2 production [9]. Taken together, these results imply that PPAR γ expression and TCR stimulation function together to regulate T cell response.

One recent study reported that pioglitazone, a PPAR γ ligand, inhibited PPAR γ -meditated differentiation of Th17, but not Th1, cells in an autoimmune EAE model, which suggested the possibility of selective regulation of Th17 cells by PPAR γ [11]. However, we found significantly increase of all the cytokines, including IFN- γ , IL-13, IL-17, and IL-9 in each T cell differentiation skewing media without affecting differentiation efficiency possibly due to hyper-proliferative property of PPAR γ deficient T cells. This increased T cell sensitivity seems to be correlated with higher AKT and ERK phosphorylation as well as rapid $I\kappa B\alpha$ degradation following TCR stimulation.

Previously PPAR γ was studied to inhibit the NF- κ B and AP-1 transcription factors in macrophages and T cells [24,25,32] and PPAR γ agonists were shown to inhibit inflammatory cytokine production in monocytes [33], as well as proliferation or IL-2 production in T cells following TCR stimulation [9]. Our results demonstrated a loss of Sirt1 and Foxo1, two negative regulators of T cell activation, in PPAR γ -deficient T cells following TCR stimulation. Sirt1, a histone deacetylase, was shown to inhibit AP-1 and is required for T cell tolerance [26]. Foxo1 is also required for inhibiting T cell activation and acts as a T cell-intrinsic regulator of tolerance [34]. The levels of Sirt1 and Foxo1 expression were significantly down-regulated in PPAR γ -deficient T cells following TCR stimulation, suggesting that PPAR γ controls the stability or expression of Sirt1 and Foxo1 to inhibit NF-kB or AP-1 activation. In addition, Foxo1 is a positive transcriptional regulator of Sirt1 and is deacetylated by Sirt1 [35]. Analysis of ChIP-seq data from BIOBASE (http://www.biobase-international.com) identified various PPAR γ response elements within the Foxo1 promoter region based on the study of genome wide analysis of PPAR γ binding region in 3T3-L1 cells [36]. Foxo1 signaling through NF-κB has been shown to mediate proinflammatory cytokine production by interacting together in a synergistic manner in macrophages [37]. Another study also revealed the role of Foxo1 in T cell tolerance by demonstrating that T cell-specific deletion of Foxo1 caused hyper-reactivity of T cells [34]. In addition, Foxo1-deleted CD4 T cells also spontaneously increased T_{FH} cells and GC formation as well as autoantibody production in 8-week-old mice [34]. ChIP analysis examining the interaction of $PPAR\gamma$ on Foxo1 promoter region will be performed in a future study to elucidate more detail mechanism.

Previous studies demonstrated treatment of T cells with a PPAR γ agonist increases Foxp3 expression and iTreg generation [10,38]. However, our results demonstrated that PPAR γ is not required for Treg generation suggesting PPAR γ agonists possibly increase Foxp3 expression PPAR γ independently. One recent study reported that PPAR γ was dispensable for Foxp3 expression in Treg cells yet critical for inhibiting colitis [39] while PPAR γ was recently shown to influence the migration of Treg cells to visceral adipose tissues by regulating chemokine receptor expression without affecting Treg differentiation [13]. Collectively, these results indicate that PPAR γ is required for the *in vivo* suppressive function related to the migration of Treg cells into inflammatory regions, but not Foxp3 expression itself.

This study is the first to characterize CD4-PPAR $\!\gamma^{\rm KO}$ mice to demonstrate spontaneous autoimmune phenotypes along with T_{FH} cells and GC formation. T_{FH} cells have recently been emerging as a T helper subset expressing Bcl-6 and IL-21 that interacts with B cells during GC formation. CD28- or ICOSdeficient mice exhibit defective GC formation, affinity maturation, and T_{FH} cell development [21,40–43]. CD28 or ICOS costimulatory signals and TCR signals are essential for the induction of T_{FH} cells [40,41]. Reports have also shown that stronger binding to major histocompatibility complex (MHC) class II and TCR is important for T_{FH} cell development [44]. Therefore, increased AKT phosphorylation and reduced expression of negative regulators to NF-KB in PPAR y-deficient T cells following TCR stimulation could contribute to increased T_{FH} cell generation in vivo. We also demonstrate an increased proportion of T_{FH} cells and number of GCs following SRBC immunization in CD4-PPAR $\gamma^{\rm KO}$ mice. In addition, PPAR γ deficient Th2 and Th17 cells produced higher IL-21 and Bcl-6 suggesting more T_{FH} cells possibly are generated and contributed GC reaction during antigen stimulation



Figure 4. Increase of CD4⁺CD62L^{low}, CD4⁺CD69⁺ and CD4⁺IL-7Ra^{high} cells in 1-year-old CD4-PPARγ^{KO} mice. Lymphocytes from spleen (SPL) and mesenteric lymph nodes (MLN) were isolated from 1-year-old control (Cre–) and CD4-PPARγ^{KO} (Cre+) mice and then stained with anti-CD4, anti-CD62L, anti-CD69 and anti-IL-7Ra antibodies for flow cytometric analysis. (A, B) CD4⁺CD62L^{low} cells, (C, D) CD4⁺CD69⁺ cells, and (E, F) CD4⁺IL-7Ra^{high} cells were analyzed. Values represent the mean \pm SEM, n=5-6. **P*<0.05. doi:10.1371/journal.pone.0099127.q004

in vivo. One previous study suggested that PPAR γ sustains Foxo1 expression [45] and another study demonstrated that mice lacking Foxo1 in T cells exhibited a large population of T_{FH} cells and a decreased number of Treg cells [25], suggesting a molecular mechanism for increase of T_{FH} cell differentiation may involving relationship of PPAR γ and Foxo1.

In conclusion, we have revealed the importance of PPAR γ in regulating T cell activation related to T_{FH} cells and autoimmunity. Modulating PPAR γ to dampen abnormal T cell activation and T_{FH} cell mediated GC reaction can be exploited for treating autoimmune diseases.

Materials and Methods

Mice

B6.129-Pparg^{tm2Rev}/J (PPARγ^{fl/fl}) mice were obtained from the Jackson Laboratory (Bar Harbor, ME). To generate CD4-specific PPARγ knockout mice, PPARγ^{fl/fl} mice were crossed with CD4-Cre^{+/-} transgenic mice. All mice were maintained at the Yale University and Hanyang University mouse facilities under pathogen-free conditions.



Figure 5. PPAR γ **deficiency in T cells results in spontaneous autoimmune phenotypes.** (A) Percentages of CD4, CD8, CD19, and NK in the spleen of 6-month-old control (Cre–) and CD4-PPAR γ^{KO} (Cre+) mice. Values represent the mean ± SEM, n = 10. **P*<0.05. (B, C) Comparison of the frequency of CD19⁺CD138⁺ plasma cells in female control (Cre–) and CD4-PPAR γ^{KO} (Cre+) splenocytes. Values represent the mean ± SEM, n = 6. **P*< 0.05. (D) The level of anti-nuclear antibody in the serum of 6-month-old mice was examined by immunofluorescence of pre-fixed NIH3T3 cells stained with diluted (1:100) sera followed by Alexa Fluor 488-conjugated anti-mouse Ig antibody. (E) Quantitation of anti-dsDNA antibody in serum of 1-year-old mice by ELISA, n = 8. (F) Hematoxylin and eosin staining of kidney from 1-year-old mice. (G) The severity of inflammation in 1-year-old mice, n = 16–19. **P*<0.05; ** *P*<0.01; *** *P*<0.001. doi:10.1371/journal.pone.0099127.q005



Figure 6. PPAR γ **deficiency in T cells spontaneously increases follicular helper T cell and germinal center formation.** (A) Representative flow cytometric gating strategy for the identification of follicular helper T cells, which were identified as CD4⁺TCR β ⁺CD62L^{low}CD44^{high}PSGL-1^{low}CXCR5⁺PD-1⁺ cells in the spleen of 1-year-old mice. (B, C) Flow cytometric analysis of T_{FH} cells from 1-year-old mice. Values represent the mean \pm SEM, n = 10. **P*<0.05. (D) Germinal centers of frozen 7-µm sections from spleen of 1-year-old mice were visualized by confocal microscopy. Slides were stained for PNA (green), IgD (red), and CD4 (blue) to detect germinal centers, B cells, and T cells, respectively. (E) The number of PNA⁺ germinal centers per spleen section was quantitated, n = 5. (F) The frequency of B220⁺GL7⁺CD95⁺ germinal center B cells from splenocytes and (G) amount of IgM and IgG1 from the sera of 1- year-old control (Cre-) and CD4-PPAR γ^{KO} (Cre+) mice was analyzed. Values represent the mean \pm SEM, n = 8–11. **P*<0.05. doi:10.1371/journal.pone.0099127.q006



Figure 7. PPAR γ **deficiency in T cells increases follicular helper T cell and germinal center formation following SRBC immunization.** (A) Experimental scheme for SRBC immunization to analyze T_{FH} cells and GC reaction. (B, C) Flow cytometric analysis of T_{FH} cells in the spleen and (D) total cell number of spleens and lymph nodes from 8-week-old control and SRBC-immunized mice. Values represent the mean ± SEM, n = 7-10. **P*<0.01, ****P*<0.01, ****P*<0.01, (E) Germinal centers of frozen 7-µm sections from spleen of 8-week-old control (non-immunized Cre-) and SRBC-immunized mice were visualized by confocal microscopy. Slides were stained for PNA (green), IgD (red), and CD4 (blue) to detect germinal centers, B cells, and T cells, respectively. (F) The number of PNA⁺ germinal centers per spleen section was quantitated n = 5-7. Expression level of (G) IL-21, (H) Bcl-6 and (I) Blimp-1 was assessed by real-time PCR and normalized to β-actin. Values represent the mean ± SEM, n = 3. ***P*<0.01, ****P*<0.01, ****P*<0.01, ****P*<0.01, ****P*<0.001, *

Ethics statement

All animal protocols used in this study were approved by the Yale Institutional Animal Care and Use Committee and Hanyang Animal Care and Use Committee.

Western blot analysis

 $CD4^+$ T cells were enriched from isolated splenocytes using antibody-coated magnetic beads. The negatively selected $CD4^+$ T cells were labeled with CD62L MACS bead (Miltenyi Biotec, Bergisch Gladbach, Germany) to purify $CD4^+CD62L^+$ naïve T cells. The purity of naïve T cells was 95~96% (data not shown). The isolated naïve T cells were stimulated with anti-CD3 and anti-CD28 antibody coated Dynabeads (Life Technologies, Carlsbad, CA) at 37°C for 0, 5, 10, and 20 min. The cells were then lysed with RIPA buffer (Cell Signaling Technology, Danvers, MA) on ice for 20 min and the protein content was determined using the Pierce BCA protein assay kit (Thermo Fisher Scientific, Waltham, MA). The proteins were resolved on 9% SDS polyacrylamide gels and transferred onto PVDF membrane (Millipore, Temecula, CA). After blocking with 5% skim milk, the membrane was incubated with specific antibodies against P-ERK, P-AKT, P-P65, IKB\alpha, Foxo1, Sirt1 and β -actin (Cell Signaling Technology) overnight at 4°C. HRP-conjugated secondary antibody followed by incubation with Supersignal West Femto ECL solution (Thermo Fisher scientific) were used to detect primary antibody binding.

Flow cytometry

Splenocytes and mesenteric lymph node cells were isolated and then stained with anti-mouse CD4-APC, CD8-PerCP-Cy5.5, CD19-PE, CD44-PE, CD62L-FITC, CD69-FITC, IL-7R α -PE, CD138-APC, PSGL-1-APC, CXCR5-PE, PD-1-FITC, GL-7-FITC, CD95-PE-Cy7, B220-Alexa647 and NK1.1-FITC (eBioscience, San Diego, CA) antibodies for 15 min at 4°C. To assess intracellular cytokine levels, cells were differentiated for 5 days and then re-stimulated with Cell Stimulation Cocktail plus Protein Transport Inhibitors reagent (eBioscience) for 5 h. Then, the cells were fixed, permeabilized, and stained with anti-mouse IFN- γ -FITC, IL-4-PE, IL-17-APC, and IL-9-APC antibodies. Intracellular Foxp3 staining was performed using the Foxp3 Staining Kit (eBioscience). To determine the purity of isolated CD4⁺CD25⁻ T cells, these cells were stained with anti-CD4 and anti-CD25, and followed by FACS analysis.

Quantitative real-time PCR

RNA from FACS-sorted CD4⁺, CD8⁺, and CD19⁺ cells was isolated using the RNeasy Mini kit (Qiagen, Venlo, Limburg) according to the manufacturer's protocol. cDNA synthesis was performed using the iScript kit (Bio-Rad, Hercules, CA). Real-time PCR was performed on a Bio-Rad CFX Connect real-time PCR detection system using iQ SYBR Green Supermix (Bio-Rad). The level of PPAR γ expression was assessed in triplicate with specific primers and normalized to β -actin. The primer sequences used were: PPARy (forward): 5'-CCAGAGTCTGCTGATCTGCG-3'; PPARy (reverse): 5'-GCCACCTCTTTGCTCTGCTC-3'; Bcl-6 (forward): 5'-CCGGCACGCTAGTGATGTT-3'; Bcl-6 (reverse): 5'-TGTCTTATGGGCTCTAAACTGCT-3'; IL-21 (forward): 5'-GGACCCTTGTCTGTCTGGTAG-3'; IL-21 (reverse): 5'-TGTGGAGCTGATAGAAGTTCAGG-3'; Blimp-1 (forward): 5'-TTCTTGTGTGTGGTATTGTCGGGACTT-3'; Blimp-1 (reverse): 5'-TTGGGGACACTCTTTGGGTAGAGTT-3'; βactin (forward): 5'-TGTCCCTGTATGCCTCTGGT-3'; and βactin (reverse): 5'-CACGCACGATTTCCCTCTC-3' (34).

ELISA

Cytokine levels in the supernatants of polarized Th1, Th2, Th17, and Th9 cultures were determined by ELISA using antibodies against mouse IL-2 (Biolegend, San Diego, CA), IFN- γ , IL-13, IL-17 and IL-9 (eBioscience) according to the manufacturer's instructions. Anti-dsDNA antibody in mouse serum was determined by ELISA (Alpha Diagnostic International Inc, San Antonio, TX).

Confocal microscopy

Mice aged 6 to 8 weeks old were immunized with sheep red blood cells (SRBC) and spleens were harvested 9 days later. Spleens from 1-year-old mice were also collected to analyze $T_{\rm FH}$ cells and GC formation. The tissues were frozen in OCT reagent and sectioned into 7-µm slices. The frozen sections were stained with anti-PNA-FITC (Sigma-Aldrich, St. Louis, MO), IgD-PE, and CD4-APC (eBioscience) overnight at 4°C. After washing, Anti-Fade reagent (Invitrogen, Life Technologies, Carlsbad, CA) was added to the slides, which were visualized using a Leica DM IRE2 confocal microscope.

T cell activation, proliferation, and differentiation

Spleens were isolated from mice and a single-cell suspension was prepared. The single cell suspension was stained against CD4 and CD25, and CD4⁺CD25⁻ T cells were purified by MoFlo (Beckman Coulter, Inc., Brea, CA), while naïve CD4 T cells were isolated using the CD4⁺CD62L⁺ T cell isolation kit (Miltenyi Biotec, Bergisch Gladbach, Germany) according to the manufacturer's instructions. Purified T cells were stimulated with platebound anti-CD3 and soluble anti-CD28 antibodies in 96-well plates and differentiated under the following subset-specific conditions: IL-12 (2 ng/ml), IL-2 (50 U/ml), and anti-IL-4 (5 µg/ml) for Th1 differentiation; IL-4 (30 ng/ml), IL-2 (50 U/

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 Chawla A, Repa JJ, Evans RM, Mangelsdorf DJ (2001) Nuclear receptors and lipid physiology: opening the X-files. Science 294: 1866–1870. ml), and anti-IFN- γ (5 µg/ml) for Th2 differentiation; TGF- β (2 ng/ml), IL-4 (30 ng/ml), and anti-IFN- γ (5 µg/ml) for Th9 differentiation; TGF- β (0.5 ng/ml), IL-6 (30 ng/ml), IL-23 (20 ng/ml), IL-1 β (20 ng/ml), anti-IFN- γ (5 µg/ml), and anti-IL-4 (5 µg/ml) for Th17 differentiation; TGF- β (5 ng/ml) and IL-2 (100 U/ml) for Treg differentiation. The cells were pulsed with H³-thymidine (1 µCi/well) overnight and H³-thymidine incorporation was measured to assess T cell proliferation.

Histology

Kidney tissues from 1-year-old mice were isolated, fixed, embedded in paraffin, and stained with hematoxylin and eosin. For each mouse, the level of kidney inflammation and more than 15 glomerular, tubular, or interstitial areas were evaluated and scored blindly for glomerular cellularity, leukocyte infiltration, severity of tubular lesions, mesangial matrix expansion, crescent formation, and interstitial mononuclear cell infiltration in the medulla and cortex. The severity of kidney lesions was determined by scoring each feature (from 0 to 3) and then calculating the mean of each set of scores. For example, glomerular inflammation was scored as follows: 0 = normal or no inflammatory cells; 1 = few inflammatory cells; 2 = moderate inflammation; and 3 = severe lymphocyte infiltration.

Statistical analysis

All data were analyzed statistically analyzed with the Student's t-test and Mann-Whitney test using Prism5 (GraphPad, San Diego, CA). p-values (P) less than 0.05 were considered statistically significant.

Supporting Information

Figure S1 Characterization of lymphocyte populations in CD4-PPAR γ^{KO} mice. (A-D) Thymic and (E-H) splenic CD4⁺, CD8⁺, and Foxp3⁺ populations from 6- to 8-week-old female wild type and CD4-PPAR γ^{KO} mice were analyzed by flow cytometry. Total CD4⁺ or CD8⁺ cells were gated from live cells while Foxp3⁺ cells were gated from CD4⁺ T cells. (I, J) Splenic B and NK cells, which were gated from CD4⁺ T cells of 6- to 8- weekold female littermate control (Cre-) and CD4-PPAR γ^{KO} (Cre+) mice were analyzed by flow cytometry. Values represent the mean ± SEM. **P*<0.05.

(DOCX)

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Author Contributions

Conceived and designed the experiments: HJP AB JMC. Performed the experiments: HJP DHK WJK JYK. Analyzed the data: HJP JYC AGS LKK JC AB JMC. Contributed reagents/materials/analysis tools: SSH ZT GRL. Wrote the paper: HJP AB JMC.

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